# Synopsis

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**Thesis Title: The Neural Basis for Host Shifts in *Rhagoletis pomonella***

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**Abstract:**

Host shifts are considered a key generator of insect biodiversity. For insects, adaptation to new host plants often requires changes in larval/pupal development and behavioural preference towards new hosts. Neurochemicals play key roles in both development and behaviour, and therefore provide a potential source for such synchronization. Here, we correlated life history timing, brain development, and corresponding levels of 14 neurochemicals in *Rhagoletis pomonella* (Diptera: Tephritidae), a species undergoing ecological speciation through an ongoing host shift from hawthorn to apple fruit. These races exhibit differences in pupal diapause timing as well as adult behavioural preference with respect to their hosts. This difference in behavioral preference is coupled with differences in neurophysiological response to host volatiles. We found that apple race pupae exhibited adult brain morphogenesis three weeks faster after an identical simulated winter than the hawthorn race, which correlated with significantly lower titres of several neurochemicals. In some cases, particularly biogenic amines, differences in titres were reflected in the mature adult stage, when host preference is exhibited. In summary, life history timing, neurochemical titre, and brain development can be coupled in this speciating system, providing new hypotheses for the origins of new species through host shifts.

**Introduction:**

Adaptation to environmental and ecological factors has been shown to play an important role in population divergence and speciation in a large number of systems (1–3). For phytophagous insects shifting to new host plants, populations potentially need to adapt their growth and development to the new host, and at the same time modulate their behavioural preference to locate that host. Furthermore, populations on novel hosts must regulate their life-history timing to coincide with the new host phenology (4–11). How these multiple events are synchronized between ancestral and novel hosts with vastly different phenologies and characteristics is an area of intense study for understanding the genesis of insect biodiversity (7,12).

*Rhagoletis pomonella* (Diptera: Tephrtidae) provides a unique opportunity to identify associations between pupal development, life history timing, and adult host choice. This species of flies originally infested fruits of the native downy hawthorn (*Crataegus mollis*) until apples (*Malus pumila*) were introduced in Eastern USA around 180 years ago (12–14). This introduction of a new host facilitated a shift in host preference from their native host, downy hawthorn, to domesticated apples, eventually leading to two host races specific to each fruit (12,13). This host shift from hawthorn to apple fruit was facilitated by two factors. First, apple and hawthorn flies are univoltine and the adults are short lived, thus each host race must emerge synchronized with the fruiting time of their host plant. Due to the earlier fruiting time of apples, apple flies initiate their overwintering dormancy, termed diapause, earlier than hawthorn flies, and emerge as adults about one month earlier as well (13). Previous work shows that the earlier seasonal adult emergence of apple host-race flies is driven solely by the timing of the termination of pupal diapause (15). The difference in adult emergence timing that drives synchronization with host fruits results in a degree of mating isolation between the apple and hawthorn races (16). Second, adults of the two host races also exhibit distinct preferences for the volatiles of their respective host fruits, which serves as an important reproductive barrier because the flies mate directly on or near the ripe host fruit (17–19). Previous studies forcing hybridisation between the host races in the laboratory showed that F1 hybrids of these two populations have altered peripheral olfactory physiology suggesting developmental

abnormalities regarding their response to the host volatiles.(20). Recent studies have shown that the change in host fruit olfactory preference between the two host races does not occur at the chemoreception stage in the antenna (21–23), but rather at the first synapse of the olfactory system in the brain, the antennal lobe (24). The coupled difference in pupal diapause timing, adult host preference, and adult brain physiology in *R. pomonella* provides a unique opportunity to examine if there are corresponding differences in brain development between the races that could simultaneously impact both developmental rate and adult olfactory host choice.

Species from temperate regions like *R. pomonella* frequently use the timing of diapause to avoid the stresses of winter and also to synchronize themselves with the phenology of their hosts the next growing season (25). While diapause is often conceived of as a state of developmental arrest, it is in fact a dynamic, physiologically regulated process with defined phases of development including the diapause preparatory stage, diapause induction, diapause maintenance, and the resumption of rapid development at the end of diapause (26,27). In insects, the central nervous system (CNS) and associated endocrine glands produce neurochemicals like neurotransmitters, neurohormones, and neuropeptides that regulate diverse physiological events including the induction and termination of diapause (28). Diapause can be regulated via changes in hormone/neurotransmitter titres, receptor abundance, or regulation of specific neurochemical pathways across the stages of diapause, such as has been found with the dopamine and serotonin pathways(29–32). Apart from life history timing, many of these same neurochemicals also play important roles in insect behaviour. Biogenic amines like dopamine, octopamine, and serotonin are known to have profound impacts on adult insect behaviour across many taxa (33–36).

The dual role of neuromodulators in regulating both insect development and host-seeking behavior suggests that changes in neuromodulator titres or production at specific life stages could impact both life history timing and preference of a phytophagous insect for its host plant through changes in brain development or differentiation. Further, the aforementioned changes in neurophysiological response and preference for host volatiles between *R. pomonella* races might be rooted in changes in brain development, neuromodulation, or both. In this study, we use a variety of chemoanalytical, morphological, and immunohistochemical techniques to examine pupal-adult brain development, life history timing, and corresponding neurotransmitter levels in two closely related populations of *R. pomonella* that differ both in diapause timing and adult preference for their respective hosts. There are a number of potential neurochemical candidates and life history stages that could play a role in this host shift, and

also a general lack of knowledge regarding how neuromodulators could impact this process. Therefore, our goal is to track brain development from larval to adult stages in the apple and hawthorn host races of *R. pomonella*, and identify which stages, if any, exhibit differences in neural development or neuromodulation between the host races, and how these differences are reflected in the mature adult fly, when host preference is exhibited.

**Methods:**

**Insect collection and maintenance**

Apple and hawthorn fruits naturally infested with larvae were collected from four different sites in Michigan, USA (Grant, Fennville, Cassopolis, Lansing) in August and September 2016, and flies were reared from larvae to adulthood following previously established *Rhagoletis* husbandry methods (14). In May to August 2017, after leaving the pupae at room temperature for 15 days, they were shipped to India (with permit). This set of pupae were used to study brain development and quantification of neurotransmitters from adult flies. Eclosed adults were maintained on a diet of sugar and yeast on a 14L:10D light cycle at 25° C and 65% humidity. Post-eclosion, young flies 1-6 days old were classified as sexually immature whereas flies that were 12-14 days old were classified as sexually mature (37–39).

To study the pre-winter and post-winter brain development as well as quantify neurotransmitters, a second set of pupae were collected in summer 2018. After collecting infested fruits from the above field sites, fruit were transferred to a tray with a wire mesh attached and kept in an insect-rearing room at 25±1°C, 14L:10D light cycle. Every day newly emerged pupae were collected and transferred to petri dishes with damp vermiculite, and maintained within a chamber containing a saturated potassium chloride solution to maintain ~85% relative humidity. To differentiate diapausing and non-diapausing pupae during the diapause initiation stage, four different cohorts of pupae were set aside and subjected to metabolic rate measurements once they reached either 7 or 19 days after pupariation. Other cohorts of pupae at 10 days after pupariation were transferred to a dark refrigerator at 4$^{o}$C with saturated KCL solution to stimulate over-wintering diapause for six months and study post winter development until they were hand-carried to India in November 2018.

**Chemicals and Reagents**

16% Paraformaldehyde EM grade (15710) was obtained from Electron Microscopy Sciences, Hatfield, PA, USA. Triton x and Bovine serum albumin were purchased from Sigma-Aldrich (Bangalore, India). All standards, ammonium acetate, Acetone, hydrochloric acid (HCl), boric acid, and reagents required for 6-aminoquinolyl-N-hydroxysuccinimidyl carbamate (AQC)

synthesis, were obtained from Sigma-Aldrich (Bangalore, India). Deuterated internal standards 14 were supplied by CDN isotope (Quebec, Canada). Ascorbic acid was obtained from Himedia (Bangalore, India), and Formic acid (FA) was obtained from Fisher Scientific (Bangalore, India). Reverse-phase solid phase extraction (RP-SPE) cartridges (Strata-X, 8B-S100-TAK) were obtained from Phenomenex, Inc. (Hyderabad, India). High-purity MS grade solvents (methanol, acetonitrile, and water) were obtained from Merck Millipore (Merck Millipore India Pvt. Ltd., Bangalore).

**Pre-diapause metabolic rate measurement**

Even though pupal diapause is ecologically obligate in *R. pomonella*, a small number of pupae avert diapause under laboratory conditions and directly develop into pharate adults in the prewinter period (11). To eliminate these non-diapausing individuals from our sampling before overwintering, we used a protocol adapted from Ragland *et* al 2009 & Powell *et* al 2021 to phenotype pupae as diapausing or non-diapausing by measuring g metabolic rates in the pre-winter period.

First we collected 7-day and 19-day-old pupae to measure their weight on an analytical balance with 5 μg precision (Mettler XP6, Toledo, OH, USA). Pupae were transferred to a 5 ml syringe used as a respirometry chamber for checking metabolic rate as an indicator of diapause or non-diapause status, held for 24h so adequate $CO_2$ could build up in the chamber, and $CO_2$ was measured on the 8th and 20th day respectively. Syringes were sealed with the plunger drawn back to produce a chamber of 1 ml internal volume. *R. pomonella* pupae were small enough to fit into one arm of the luer valve, allowing the full volume of the syringe to be injected. We also purged multiple empty syringes to serve as controls/blanks. Every 24 h, the full volume of each syringe was injected into a flow-through respirometry system consisting of a Li-Cor 7000 infrared $CO_2$ analyser (Lincoln, NE, USA) with a resolution of 5 parts per million (ppm) $CO_2$ interfaced to a Sable Systems International UI-2, recorded by Expedata data logging software (Las Vegas, NV, USA). The flow rate was fixed at 150 ml/min using a Sierra Instruments mass flow controller (Monterey, CA, USA). $CO_2$-free air, scrubbed with a dririte-ascarite-dririte column, served as the baseline for measurements, and the system was routinely calibrated with $CO_2$-free air and a certified standard mixture of 500 ppm $CO_2$ in nitrogen (Airgas, Jacksonville, FL, USA). For one replicate a total of 30-40 pupae were used to calculate the metabolic rate and after that, the brain samples were dissected on cold PBS(1x) and flash frozen with liquid nitrogen. The total number of brains dissected are denoted as $\mathbf{n_{dissections}}$, and total number of

brains used for staining denoted as $\mathbf{n}_{staining}$ for each sample. For further details, see Supplementary Materials, Extended Methods.

**Brain morphology and immunohistochemistry**

In early November 2018, diapausing pupae were hand-carried with ice packs from Gainesville, FL, USA to Bangalore, India. After keeping pupae 6-months in the refrigerator (October 2018 - March 2019), in March 2019 all four different cohorts of pupae of the apple race were pulled out at their respective 6-month time points and left at room temperature 25°C, 14L:10D light cycle, and 65% humidity as described in Insect collection and maintenance above. Every 5 days after removal from artificial overwintering, a few pupal caps (n=5-35) were removed to observe development externally and brains were also dissected from each individual to assess brain development (Figs. 1-2). Further, pupae were individually photographed using an infinity HD camera (lumenera, model number N9033210) attached to a stereomicroscope. After that, the brain was individually dissected using 50ul of 1X phosphate buffered saline solution, PBS (PH: 7.4) on ice with fine forceps, and stored at -80$^{o}$C. Four different cohorts of pupae of the Hawthorn race were pulled out in April 2019 at their respective 6 months of overwintering time points (November 2018 - April 2019) and were photographed and then dissected using the above protocol.

To characterize brain morphology, brain samples were dissected as mentioned above and then underwent immunohistochemical staining using a protocol adapted from those used in *Drosophila and Rhagoletis* (24,40,41). Further details can be found in Supplementary Materials, Extended Methods.

**Quantification of neurotransmitters from the R. *pomonella* brain**

After morphologically determining developmental stage, brain samples were removed from -80°C storage and five brains of the same stage were pooled for each sample replicate (denoted as $\mathbf{n}_{samples}$). In one sample, only a single brain at stage 4 was observed in the hawthorn race prior to day 35 after the artificial winter. To assess the neurochemistry of this brain, we were required to pool it with other hawthorn stage 4 brains from days 35 and 40 to obtain enough material for analysis. This unique data point is shown at day 30 in Figure 4. The sample processing was performed as in Ramesh *et* al, 2019 (42–44). After the brain samples were pulled out, samples were immediately transferred to a vial containing 190 µl of Acetone (with 0.1% Formic acid, FA) and 10 µl of 1% of ascorbic acid (1.76mg/ml), followed by derivatization with 6-aminoquinolyl –N-hydroxysuccinimidyl carbamate (AQC) as in Ramesh et al 2019. (42–44). Adult fly brains were dissected individually, pooled into a group of five,

and stored at -80ºC with190 µl of Acetone and 10 µl of 1% of ascorbic acid. After that 10 µl of internal standard (ISTD), a mixture of all 14 neurotransmitters, (0.5ug/ml i.e. 1ng on column) was added to brain samples in screw cap vials. This mixture was sonicated for 1 min, and then homogenised using a plastic pestle. It was then immediately centrifuged (13500 rpm 4ºC, 5 minutes), and the supernatant was transferred to a new tube. Simultaneously, the vials for standard solutions for calibration curves were also prepared (supplementary table s11). Serial dilutions of standard stocks were prepared with highest concentration considered on column 100% to five points including 50%,25%,12.5% and 6,25% of the maximum quantity for each of the targeted compounds, for pre-winter and post-winter brain samples. For sexually immature and sexually mature adults the standard stock was diluted to prepare 200%, 160%, 80%, 40%, 20% of the targeted compounds. After that 10 µl internal standards with 190 µl acetone (0.1% FA) and 10 µl of ascorbic acid were added to the standard solution tubes. All samples and standard tubes were dried in a speed-vac for 1hr. Apart from these, one more vial containing 16 amino acids (1 µl of 10 µg/ml) was added as an additional standard. These 16 amino acid standards helped to validate the method and retention time used for mass spectrometry (MS) every time we ran a sample. Once everything was dried in the speed-vac, 80 µl of borate buffer with 10 µl of ascorbic acid (1.76 mg/ml) was added to all tubes and vortexed. Before analysis, 10 µl of 10 mg/ml AQC (prepared in 100% Acetonitrile, ACN) was added and kept for 10 min at 55ºC. After that 3ul of 100% formic acid FA was added and the tube was vortexed to stop the reaction. Tubes were subsequently kept at room temperature until all SPE columns were rinsed and cleaned with 100% methanol and 0.1% FA. After that, 500 µl of water was added to all the tubes, then vortexed and the solution was loaded on SPE columns. Columns were washed twice with 0.1% formic acid prepared with LC grade water. After that 1ml of ACN: MeOH 4:1 in 0.1% formic acid was added to the column and eluted in a new vial. All tubes were dried in a speed vac for 3hrs. Dried samples were stored at -20ºC until they were run on LC-MS. Each sample was thawed and reconstituted in 50 µl of 2% ACN prepared in 0.5% FA. The LC-MS instrument method and setup are described in detail in tables s9, s10, s11, s12 figure s6 and s7. The final quantification and analyses can be found in detail in Supplementary Materials, Extended Methods.

**Statistical Analysis:**

1) To compare the relative differences in timing of the proportion of apple vs. haw flies transitioning from pupal diapausing brain morphology to adult brain development post winter, we used a generalised linear model with a binomial, log-link function in R (V4.0.2) with ggplot,

dplyr, and tidyr packed. In this logistic GLM host, days after winter was used to explain the proportion of individuals transitioning from stage 3 to stage 4 of brain development between apple and hawthorn flies, as well as to estimate 95% CIs around the logistic estimates. 2) Respiration and neuromodulator data were analysed using linear mixed effects models in R (V4.0.2) using the lmer function from the lme4 package. In the models, age and host race were used as a fixed effect whereas cohort / batch number was used as a random effect. Further, Tukey's HSD tests with correction for multiple comparisons were performed with pairwise multiple comparisons using the lmerTest package. Satterthwaite approximations on the lmerTest package were used to test the significance of the effects. 3) The interaction between host race and days on amount of precursor neurochemicals at brain development stage 4 was done using, generalised linear model, Univariate analysis of variance in SPSS V.26. In the model host race and days were used as fixed factors and quantities of each neurochemical was used as the dependent variable. Further, Tukey's LSD tests were performed for pairwise comparisons with significance cut offs of 0.05.

**Results:**

***Rhagoletis pomonella* host races exhibit distinct brain development stages during pupation**

We observed external morphological and brain development in *Rhagoletis pomonella* pupae during diapause initiation, pre-winter diapause, post-winter diapause, diapause termination, pharate-adult, and eclosed adult developmental phases (Figure 1 and s2). The entire set of experiments were performed over a period of two seasons 2017 to 2019 with total of 2,091 individuals examined across both the host races. These phases were divided into 8 different stages according to changes in external head morphology, brain development, or metabolic rate (Figure 1, s1 and s2). The pre-winter diapause initiation stage 1, (8 days after puparium formation, Methods & Figure s1, and s2 apple $\mathbf{n}_{\mathbf{dissections}} = 75$, $\mathbf{n}_{staining} = 7$, hawthorn $\mathbf{n}_{\mathbf{dissections}} = 65$, $\mathbf{n}_{staining} = 8$), and the pre-winter diapause maintenance stage 2, (20 days after puparium formation, Figure s1, s2, Figure 1, apple $\mathbf{n}_{\mathbf{dissections}} = 65$, $\mathbf{n}_{staining} = 7$, hawthorn $\mathbf{n}_{\mathbf{dissections}} = 75$, $\mathbf{n}_{staining} = 8$) were differentiated using respiration rates. Wherein pupae entering diapause exhibited higher metabolic rates 8 days after pupariation that stabilized at low levels of metabolic depression by 20 days after pupariation, when pupae were clearly in the diapause maintenance phase [Figure s1,(15,45)]. During these two pre-winter stages, brain morphology reflected that of the larval brain with a clearly identifiable ring gland and sub-oesophageal ganglion (Figure s2, A). After a six-month artificial overwintering period at 4-5$^{o}$C, we observed

substantial neural differentiation and remodelling had occurred compared to pre-winter diapausing brains (stage 3, Figure s2, B, apple $\mathbf{n_{dissections}}$ = 134, $\mathbf{n}_{staining}$ = 2, hawthorn $\mathbf{n_{dissections}}$ = 171, $\mathbf{n}_{staining}$ = 3) in that the ring gland became disassociated and the place between the hemispheres containing the dorsal vessel became thin (c.f. stages 1 & 2; Figure s2, A, B & movie 2, 3).

Subsequent assessment of later stages also showed that adult brain (CNS) development and differentiation occurred after overwintering, at a stage during pharate-adult development where early antennal development and changes in head morphology were also observable (stage 4, Figure s2 C, apple $\mathbf{n_{dissections}}$ = 70, $\mathbf{n}_{staining}$ = 4, hawthorn $\mathbf{n_{dissections}}$ = 60, $\mathbf{n}_{staining}$ = 1). At this stage 4, brain region boundaries were apparent, but the regions themselves remained undefined (Figure s2, C). By stage 5, orange pigmentation had accumulated in the eyes and a transparent antenna became well developed. At this stage, the central nervous system was more defined with structures including the antennal lobe, mushroom body, and suboesophageal ganglion (Figure s2, apple $\mathbf{n_{dissections}}$ = 43, $\mathbf{n}_{staining}$ =1, hawthorn $\mathbf{n_{dissections}}$ = 35, $\mathbf{n}_{staining}$ = 2). The last three stages (including the sexually immature and mature eclosed adult stages) were similar in both brain and external head morphology, just with greater progression of bristle development and pigmentation of the pharate-adult cuticle (Stage 6, apple $\mathbf{n_{dissections}}$ = 20, $\mathbf{n}_{staining}$ = 1, hawthorn $\mathbf{n_{dissections}}$ = 41, $\mathbf{n}_{staining}$ = 1; stage 7, apple race $\mathbf{n_{dissections}}$ = 45, hawthorn race $\mathbf{n_{dissections}}$ = 50, and stage 8 apple race $\mathbf{n_{dissections}}$ = 45, hawthorn race $\mathbf{n_{dissections}}$ = 45, hawthorn $\mathbf{n}_{staining}$ = 1 Figure s2, D).

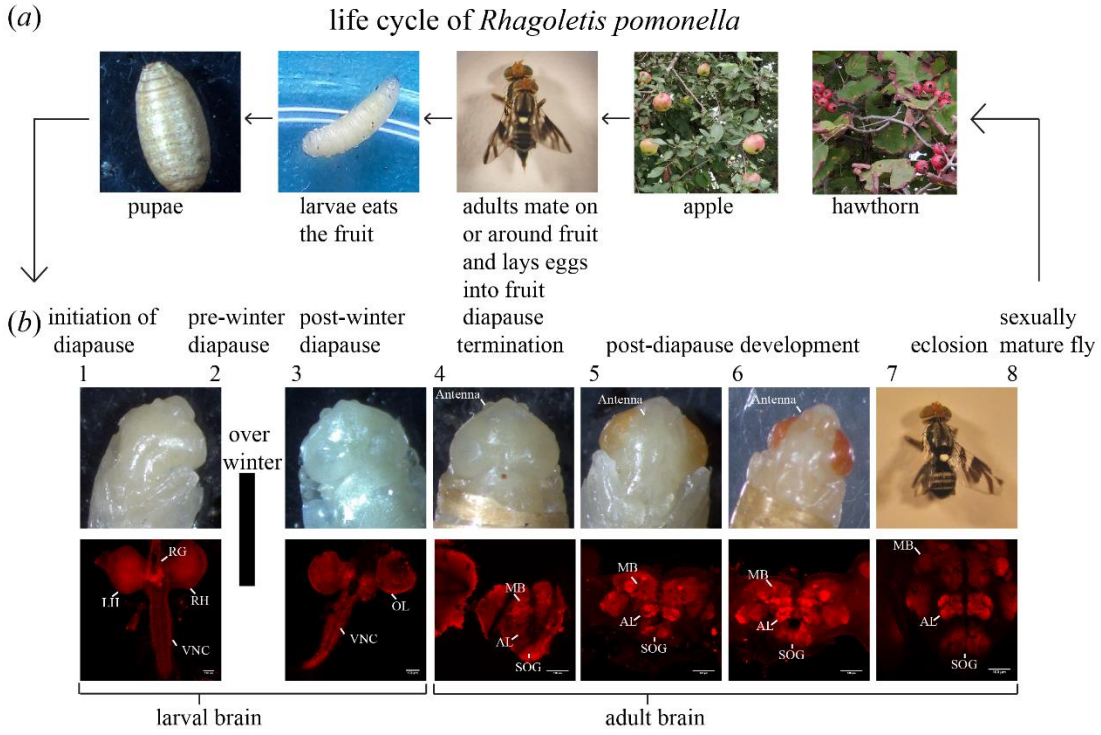


**Figure 1:** The life cycle of *Rhagoletis pomonella.* (A) Life stages of the fly from mating to oviposition from right to left. (B) Micrographic images of pupal, pharate-adult, and adult developmental stages from left to right. The upper stereomicrographs in the bottom panels show the head morphology while the lower confocal micrographs indicate the corresponding brain morphology of those stages using immunohistochemical nc82 staining across the different developmental stages (stages 1-8). The developmental stages are classified based on distinct head morphology, metabolic rate, or CNS development. Our imaging data showed that there are no morphological differences between stages 1 and 2 or between stages 7 and 8. The stages are as follows: **1**, Pre-winter diapause induction stage pupae still have high metabolic rates (Methods & Figure s1, s2, apple race, stage 1); **2**, pre-winter diapause induction stage when pupae have entered metabolic depression (Figure s1, s2 Figure 1, hawthorn race, stage 2); **3**, post-winter diapause maintenance phase (Figure s2, apple race, stage 3); **4**, end of diapause (Figure s2, hawthorn race, stage 4); **5**, midway through pharate-adult development (Figure s2, apple race, stage 5); **6**, late pharate-adult development ((Figure s2, apple race, stage 6); **7**, sexually immature adult fly, less than 7 days old ((Figure s2, hawthorn race, stage 7); **8**, sexually mature fly, more than 12 days old; The anatomical brain regions are identified as left and right brain hemispheres (LH and RH), ring gland (RG), optic lobe (OL), ventral nerve code (VNC), the antennal lobes (AL), mushroom bodies (MB), and suboesophageal ganglion (SOG).

**Apple race pupae exhibit more rapid onset of adult brain morphogenesis than the hawthorn race**

To compare post-winter brain development between the two races of *R. pomonella*, both apple and hawthorn pupae were brought to room temperature (25$^{o}$C) after 6 months of simulated overwintering (4-5$^{o}$C) synchronized to their respective diapause initiation timing (Figure 2 A, B and Tables s1 and s2). By sampling a subset of pupae every 5 days from day 0 to day 70 after removal from artificial overwintering conditions, we observed that both host races exhibited onset of adult neurogenesis during the post-winter phases, as shown in Figure 1. In the apple race, a substantial number of pupae exhibited adult brain morphology (Stage 4-6 in Figure 1) starting from day 0 after removal from simulated winter and until day 25 when 42% of the pupae sampled exhibited adult brain morphology (57/135). Conversely, in the hawthorn race only 4/152 (3%) total pupae exhibited adult brain morphology even until day 40 (Figure 2 A, B and Tables s1, s2). Logistic regression analysis (Figure 2 C) showed that apple race individuals began initiating adult brain development significantly more rapid than hawthorn race individuals ($x^2$day = 265.79, $p<0.001$; $x^2$host = 88.04, $p<0.001$; $x^2$day*host = 34.2, $p<0.001$). Diapausing apple race pupae began adult brain development approximately 24 days faster than diapausing hawthorn race pupae, with 18.34 ($\pm$0.56-0.42,95%CI) days for 50% pupae exhibiting adult brains in apple vs. 42.10 ($\pm$0.61-0.42,95%CI) days for hawthorn (Figure 2). Therefore, even though the artificial overwintering period was of the same duration for both the races, the transition from pupal to adult brain development began occurring significantly more rapid in the apple race vs. the hawthorn race. In other words, not only has the apple race shifted its overall life cycle to coincide with host phenology, but the rate of development of the pupal brain from the post winter to the initiation of adult brain morphogenesis has become more rapid.

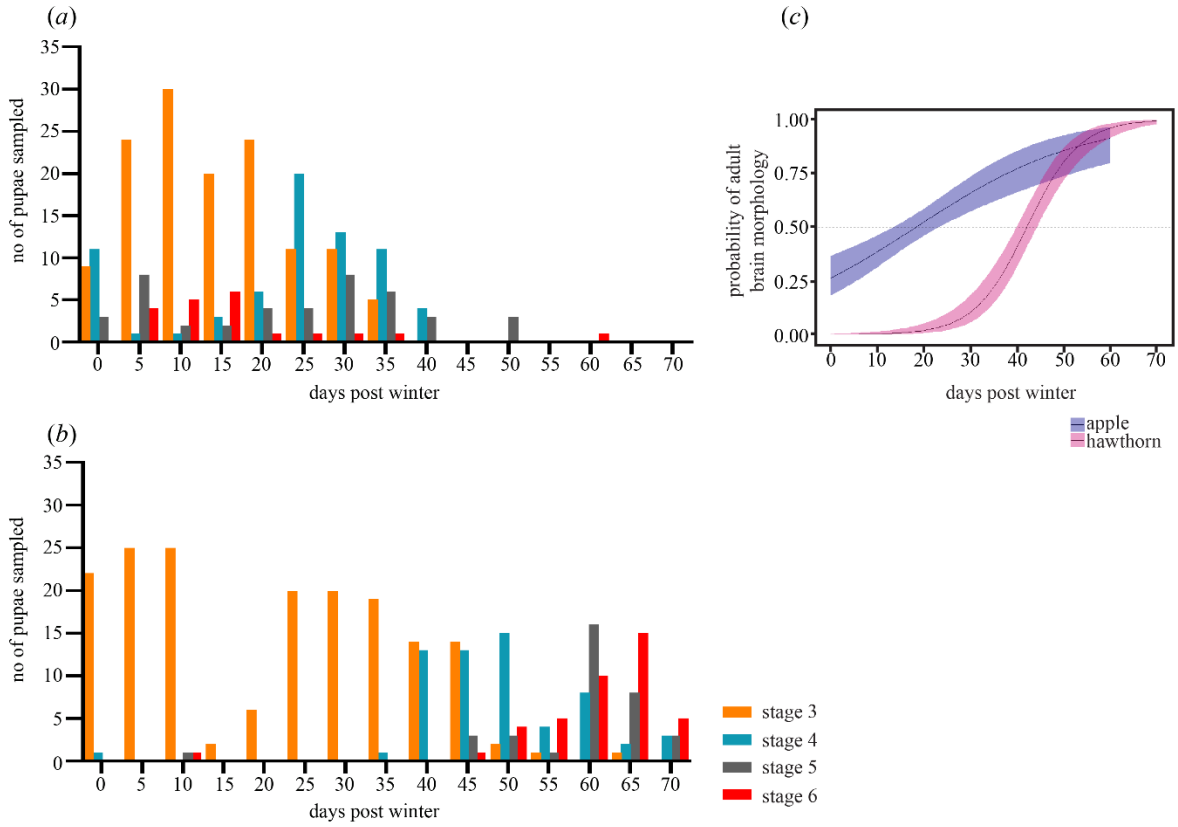


**Figure 2:** Post-winter brain development over time between the two host races. Brains were dissected from pupae and assessed every 5 days after simulated overwintering. A) Apple race, stages 3, $\mathbf{n_{dissections}}$ = 134; stage 4, $\mathbf{n_{dissections}}$ = 70; stage 5, $\mathbf{n_{dissections}}$ = 43; stage 6, $\mathbf{n_{dissections}}$ = 20; B) Hawthorn race, stages 3, $\mathbf{n_{dissections}}$ = 171; stage 4, $\mathbf{n_{dissections}}$ = 60; stage 5, $\mathbf{n_{dissections}}$ = 35; stage 6, $\mathbf{n_{dissections}}$ = 41; C) Logistic regression curves with 95% confidence intervals for the proportion of adult brain morphology vs. larval brain morphology observed in *R. pomonella* apple and hawthorn race pupae over time after artificial overwintering.

**Neuromodulator levels in developing Hawthorn race pupal brains are generally higher than in the apple race.**

We next examined a total of 14 neurochemicals across 6 biochemical pathways in the developing brains of both hawthorn and apple race of *R. pomonella.* These neurochemicals included both precursor molecules and their products (Figures, 3, s3, s4 & s5). We used linear mixed models to assess both within and between host race comparisons. All statistical results have been corrected for multiple comparisons using Turkey's HSD. In general, the titres of precursor chemicals increased first in the post-diapause stages as pharate-adult brain development began (transition from stage 3 to 4), whereas product molecule levels did not change until later stages, even as late as post-eclosion adult fly sexual maturation. Of the 14 neurochemicals examined, only four chemicals showed higher titres in the apple race,

(dopamine, stage 5, p= 0.0001; serotonin, stage 6, p= 0.0001; serine, stage 6, p= 0.009). For all other molecules, the hawthorn race exhibited higher titres than the apple race. One exception is tyramine which showed higher titres in the hawthorn race at the pupal stages and higher tires at the sexually mature adult stage in the apple race (tyramine, stage 4, p= 0.0001; stage 6, p= 0.039; stage 8, p= 0.0114). These chemicals further increased in titre as *R. pomonella* development progressed, until the point of adult fly sexual maturity, when host preference first emerges. At the stage of fly sexual maturity (Stage 8, Fig. 1), product neurochemicals from two major pathways showed a difference between the host races involving histidine to histamine and tyramine to octopamine, respectively (stage 7, apple race $\mathbf{n_{samples}}$ =10, $\mathbf{n_{dissections}}$ = 50; hawthorn race $\mathbf{n_{samples}}$ =10, $\mathbf{n_{dissections}}$ = 50; histidine, p = 0.005; stage 8, apple race $\mathbf{n_{samples}}$ =10, $\mathbf{n_{dissections}}$ = 50; hawthorn race $\mathbf{n_{samples}}$ =10, $\mathbf{n_{dissections}}$ = 50; tyramine, p= 0.011; octopamine, p= 0.011; histamine, p= 0.020).

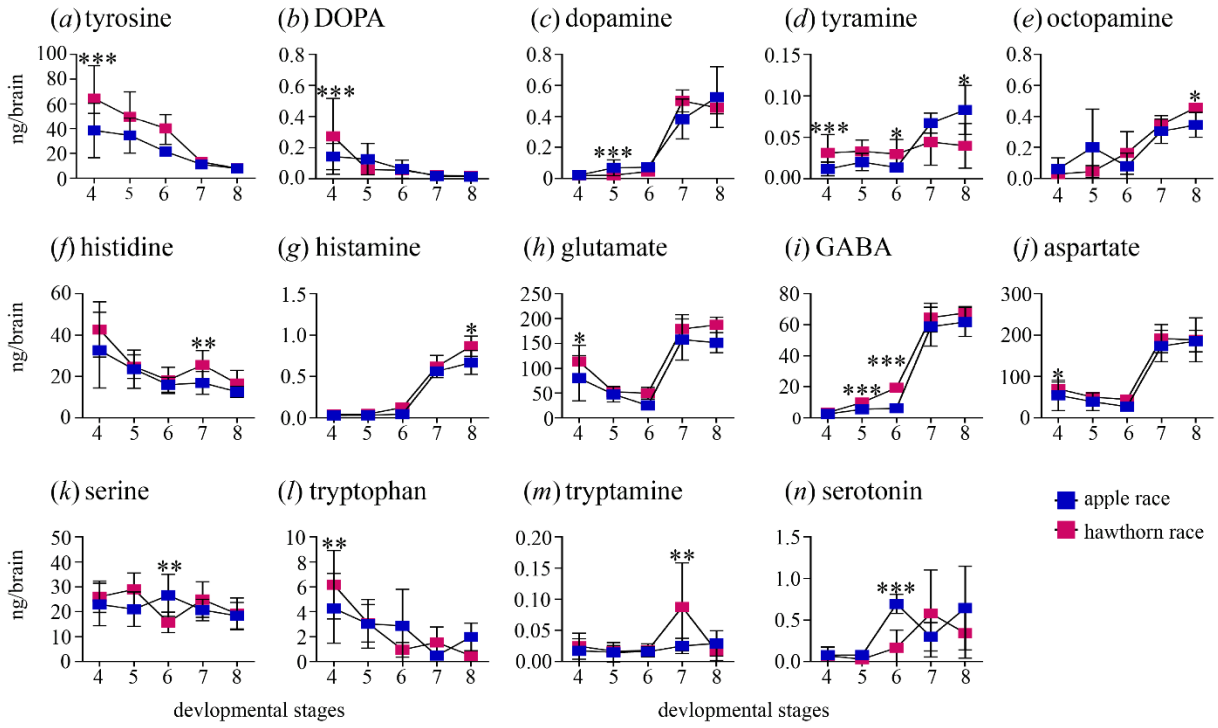


**Figure 3:** Quantification of biogenic amines and their precursors from the onset of adult brain development (stage 4) all the way up to sexually mature fly (stage 8) as defined in Figure 1 (A-N). Line graph of neurochemical titres for both host races at different developmental stages with 4-15 $\mathbf{n_{samples}}$ per stage, containing a pool of five brains in each sample, symbols represent mean with SE: A) tyrosine; (B) DOPA; (C) dopamine; (D) tyramine; (E) octopamine; (F) histidine; (G) histamine; (H) glutamate; (I) GABA; (J) aspartate; (K) serine; (L) tryptophan; (M) tryptamine; and (N) serotonin; Asterisks above indicate differences between host races at the equivalent stage of brain development. P-values

represented are < 0.05 *, < 0.01 **, and < 0.001 ***, linear mixed effect model, followed by Tukey's HSD correction for multiple comparisons.

**Rapid onset of adult brain morphogenesis in the apple race corresponds to lower levels of neuromodulators**

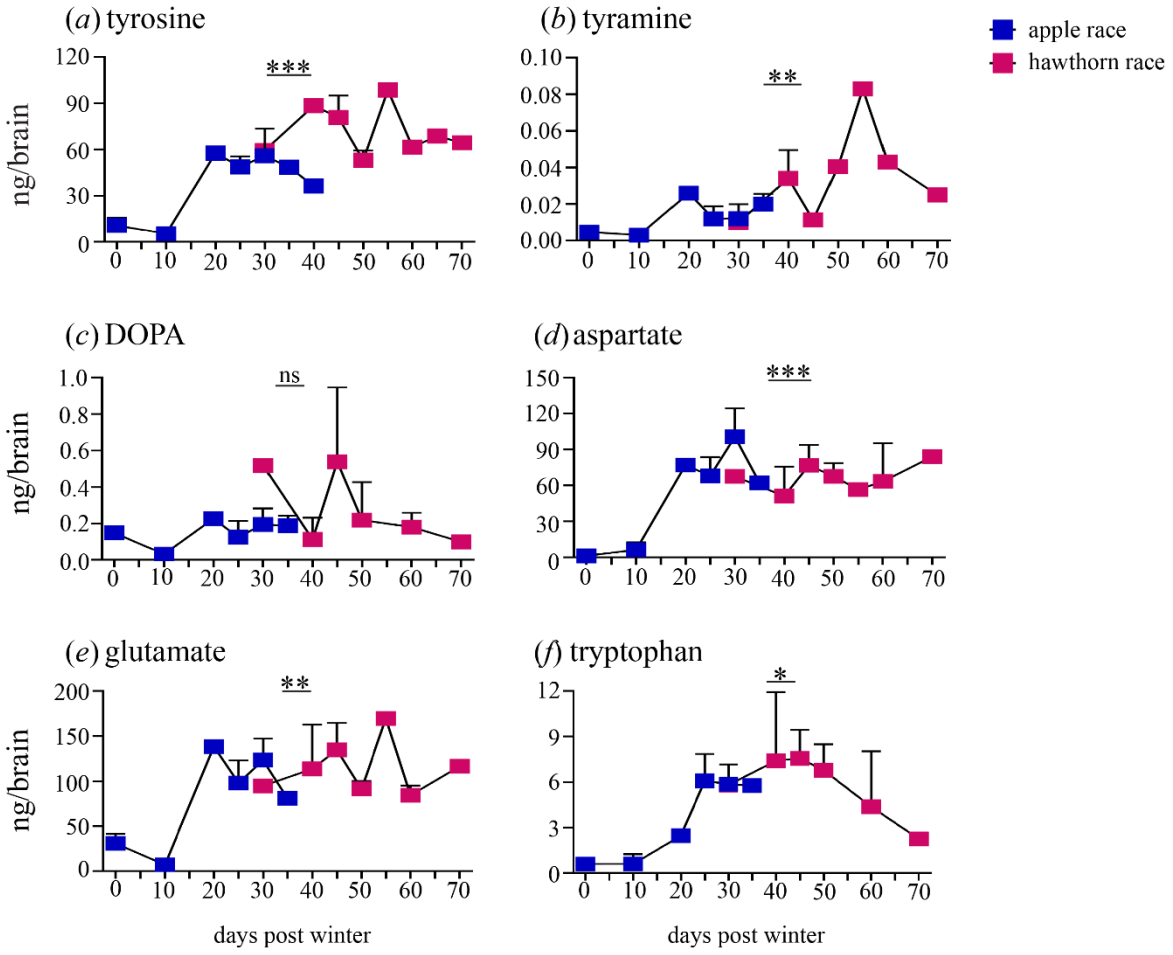


**Figure 4:** Quantification of precursor neurochemicals during onset of adult brain morphogenesis. (A-F) Line graph of neurochemical titres for both host races at different developmental stage 4 with 1-4 $\mathbf{n_{samples}}$ per time point, containing a pool of five brains in each sample, symbols represent mean with SE: A) tyrosine; B) tyramine; C) DOPA; D) aspartate; E) glutamate; F) tryptophan; Asterisks above indicate significant interactions between effects of host race*days for neurotransmitter titre. P values represented are < 0.05 *, < 0.01 **, and < 0.001 ***, analysis using Two-way ANOVA, post hoc Tukey's LSD test.

Given that in the apple race, the rate of adult brain development is significantly more rapid than the hawthorn race (Figure 2), and also exhibits lower titres of several neuromodulators, we next assessed whether the rate of development corresponded to lower levels of neuromodulators, particularly at the first stage of adult brain development (Stage 4). Figures 3, s3, s4 and s5 show that six precursor molecules including tyrosine, tyramine, DOPA, aspartate, glutamate, and

tryptophan from four different biosynthetic pathways showed significant differences in titre between the host races at the first appearance of the adult brain during pupal-pharate adult metamorphic development, stage 4 in Figure 1 (stage 4, apple race $\mathbf{n_{samples}}$ =14, $\mathbf{n_{dissections}}$ = 70; hawthorn race $\mathbf{n_{samples}}$ =11, $\mathbf{n_{dissections}}$ = 55; tyrosine, p= 0001; DOPA, p= 0.0004; tyramine, p= 0.0001; aspartate, p= 0.022; glutamate, p= 0.002; tryptophan, p= 0.002;). For each chemical, the titres were higher in the hawthorn race compared to the apple race. To compare how these precursor neurochemicals titres changed over time between the developing hawthorn and apple race pupae, we compared neurochemical titres of Stage 4 brains against the day they were sampled after winter.

A Univariate analysis of variance, UNIANOVA was conducted to examine the effects of host race and days after winter on the titres of these six precursor neurochemicals. Except for the neurochemical DOPA, there was a significant interaction between host race and day for each neurochemical titre (table s7, tyrosine, F (12,10) =10.9, P = 0.0001; tyramine, F (12, 10) = 4.95, p = 0.008; DOPA, F (12, 7) = 0.82, p = 0.632; aspartate, F (12, 11) = 10.1, p= 0.0001; glutamate, F (12, 11) = 6.47, p = 0.002; tryptophan, F (12, 10) = 4.35, p = 0.013). Therefore, neurochemical titres were significantly lower in rapidly developing apple race pupae than at later time points, when hawthorn race pupae were beginning to develop adult brain morphology. This was true for precursor chemicals even when accounting for pupae that might have been impacted by the temperature effects of development during winter itself (i.e. until day 20) (table s8, tyrosine, F (8,9) =38.14, P = 0.0001; tyramine, F (7, 10) = 4.46, p = 0.017; DOPA, F (7, 8) = 0.84, p = 0.581; aspartate, F (8, 10) = 14.25, p= 0.0001; glutamate, F (8, 9) = 25.36, p = 0.0001; tryptophan, F (7, 10) = 7.19, p = 0.003). As a result, more rapidly developing brains in apple race pupae exhibited significantly lower titres of neuromodulators (especially biogenic amine precursor molecules) than later developing hawthorn race brains, and these differences were also reflected in product molecules at the adult fly stage at which host preference is exhibited (compare stages 4 and 8 in Figures 3, s3 and s5). (37–39).

**Discussion:**

In this study, we have identified differences in rates of adult brain morphogenesis and levels of several neurochemicals between the apple and hawthorn host races of *Rhagoletis pomonella* across the transition from pupal diapause to post-diapause, pharate-adult development, and in the adult fly. These two closely related populations differ in both life history timing, adult host preference and are generally considered a model for incipient ecological speciation. The divergence of these two populations towards hosts with such different phenologies serves as

an excellent system to examine how multiple life history events are synchronized between ancestral and novel hosts.

First, we found that while some neurogenesis occurred during the diapause maintenance phase in both host races, substantial adult brain development and differentiation was initiated only after winter and brain development progressed through several morphological stages to adult emergence. In particular, pre-diapause individuals were identical in both host race and neurochemical titres (stages 1 and 2). This indicates that any difference between the host races occurs only after diapause, and is in agreement with recent transcriptomic work showing rapid up-regulation of growth and development-related transcript genes in the apple race with shorter post-winter diapause duration as compared to the longer post-winter diapause duration of hawthorn race (46). When we dissected brains from apple and hawthorn race pupae at regular intervals after they were removed from overwintering, almost no hawthorn pupae showed brain development beyond stage 3 for at least a month after overwintering, and it took up to 50 days for the majority of hawthorn pupae to terminate diapause and transition from stage 3 to stage 4 where pharate adult neural development was apparent (Figure 2B). In contrast, some apple race pupae were already progressing to stage 4 pharate adult development as soon as they were removed from overwintering (Figure 2B). This finding indicates not only has the apple race shifted its entire life cycle to coincide with fruit phenology, but adult brain morphogenesis itself occurs roughly three weeks faster in the apple race as compared to the hawthorn race even when the overwintering period is synchronised between the host races. This change in adult brain morphogenesis is not necessarily predicted from its overall host shift, and suggests a unique phenotype in the derived apple host race.

To better understand how neurochemical signaling could be associated with the different stages of development in *R. pomonella*, we examined six major biosynthetic pathways known to impact brain development as well as behavior in insects (42,47). Our results indicate that the titres of 11 out of 14 neurochemicals were significantly reduced in the apple race pupae across multiple developmental stages as compared to the hawthorn race (Figure 3, e.g., tyrosine, tyramine, octopamine, DOPA, histidine, histamine, aspartate, glutamate, GABA, tryptophan and tryptamine). Further analysis showed that these lower titres also corresponded with earlier development of the adult brain in the apple race post diapause (Figure 4, e.g., tyrosine, tyramine, aspartate, glutamate, tryptophan).

Our main goal of this study was to link life history timing, brain development and neurochemistry in the *Rhagoletis* system by identifying particular developmental stages and

neurochemicals that differ between the host races. Here, we have identified specific differences in several neuromodulators, particularly the biogenic amine pathways for octopamine and dopamine, at the first appearance of the adult brain (stage 4) and again at sexual maturity in the adult fly, when host preference is exhibited (stage 8). These differences are further coupled with a difference in developmental timing of adult brain morphogenesis between the two races. This data therefore provides specific timepoints and chemical pathways that could be assessed to determine if and how they impact life history and physiology in these populations.

For example, we show that onset of adult brain differentiation (Stage 4) also corresponds with morphogenesis and emergence of the adult antenna, and this stage is accompanied by several significant differences in neuromodulator levels between the host races. A recent study comparing olfactory neurophysiology between the apple and hawthorn races of *R. pomonella* identified a neuronal switch in the chemosensory system in the adult brain associated with differential host choice behaviour towards apple or hawthorn fruit (24). Such a switch in neurophysiology has similarly been observed between the Z and E strains of the European Corn Borer, *Ostrinia nubilalis*, where male preference for a particular isomer of the sex pheromone is controlled by cis-acting variation in a sex-linked transcription factor (bric à brac; bab) expressed in the developing male antenna (48). A recent study of the *Rhagoletis* brain transcriptome also shows variation in cis-regulatory elements associated with differentially expressed transcripts during diapause development and an important role of hub genes in transcriptional networks that differ during diapause development between the two host races (46). Interestingly, in *Drosophila* cis-regulatory variation in two genes involved in these pathways, tyrosine hydroxylase and dopa decarboxylase (Figure 3A), have been shown to impact neurogenesis (47). Future studies that measure the expression of enzymes involved in the production of dopamine and octopamine and selective pharmacological treatments that act as agonists and antagonists of these biogenic amines at early developmental stages in these two host races could indicate if these pathways are involved in the differentiation of these two host races.

To conclude, we have characterized the progression of neurogenesis from diapause onset to adult reproductive maturation in both the apple and hawthorn host races of *R. pomonella*, an important model for ecological speciation and diversification. We identified significantly lower neurochemical levels, particularly biogenic amines in the dopamine and octopamine pathways, in the apple race of *R. pomonella* that correspond to more rapid adult brain morphogenesis in this new host race. These differences in neurochemical titre between the races in the developing

pupal stages are also apparent in the adult brains at stages when flies are reproductively mature. Because biogenic amines have been implicated to impact both pupal diapause and adult behaviour, this study offers a new hypothesis that could correlate life history timing and adult host preference through developmental differences in neuromodulation. This hypothesis must now be tested in further studies assessing enzymatic expression and pharmacological manipulation of neuromodulator levels in developing pupae. As previously suggested, connecting host preference and survival through relatively simple changes could be a widespread mechanism for generating biodiversity across phytophagous insects, contributing to the origin of the large number of species observed (24).

**Data accessibility:** All data supporting this manuscript will be uploaded to Dryad upon submission.

**Authors' contribution:** HK, SBO, and DAH conceived the study and designed experiments; JLF provided the adult flies and reagents for the experiment. HK conducted experiments; HK, DAH and SBO analyzed data; HK and SBO wrote the manuscript; all authors revised and approved the manuscript.

**Competing interests:** The authors declare no competing interests.

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**Supplementary Information:**

**Supplementary Data File:** This file contains extended methods, figures s1-s7, tables s1-s12.

**Extended Methods:**

**Pre-diapause metabolic rate measurement**

Previous work had shown that metabolic rates dropped as pupae initiate diapause and become stabilized after they enter into the diapause maintenance phase, with a fraction of pupae not entering diapause and developing directly into pharate adults that are identifiable by their much higher metabolic rates (1). We initially measured metabolic rates and dissected individual pupae on day 6 (n=70), day 8, day 15 (n=40), and day 20 after pupariation. This initial effort showed that day 6 individuals were still undergoing larval-pupal metamorphosis and by day 8 they showed fully developed pupal morphology. We therefore measured individuals on day 8 after pupariation to estimate metabolic rates during diapause initiation. By day 20 after pupariation, metabolic rates were lower than on day 8, showing that pupae had transitioned from diapause initiation and entered into the metabolic depression indicative of the diapause maintenance phase (Figure s1) (1,2). Because we wished to conservatively remove any individuals that may not have been in diapause from the experiment, any individual with a metabolic rate higher than 0.025 ul $CO_2$/mg/hour was eliminated from further sampling. When we dissected pupae in the low metabolic rate class indicative of being in the diapause maintenance phase at day 20 after pupariation, they clearly possessed the larval brain morphology seen in stage 2 (Fig. 1). However, when we dissected brains from pupae with metabolic rates higher than 0.025 ul $CO_2$/mg/hour 20 days after pupariation these brains were in stage 4 (Fig. 1), indicating pharate-adult development had begun.

**Brain morphology and immunohistochemistry**

Briefly, dissected brains were transferred into plastic cell culture plates containing 4% paraformaldehyde (PFA), and plates were covered with aluminium foil. Brain samples were fixed in PFA for 3 hours on a rotator at 4°C under red light and subsequently washed with 0.3% Triton X prepared in 1x PBS (PTX) solution three times at an interval of 15 minutes. Brain samples were then treated with 0.1% bovine serum albumin in PTX (PBTX) as a blocking agent for 15 minutes, followed by incubation with the primary mouse anti-Bruchpilot/mAbnc82 (1:30, DSHB, University of Iowa) antibody for 48 hours on a rotator at 4°C. After two days of incubation with antibody, samples were washed four times with PTX solution at 15-minute intervals. Secondary antibody goat anti-mouse Alexa 647 (1:400, Invitrogen, Thermo Fisher Scientific, Waltham, USA) was added and brain samples were

incubated for 48 hours on a rotator at 4°C. Mounting of the brain for visualization occurred on the fifth day when samples were washed with PTX 4 times, at an interval of 15 minutes at 4°C, and then mounted in 70% Glycerol on glass slides. Optical sections (512 x 512 pixels) of each brain sample with 1 µm step size, were imaged with a confocal scanning microscope (Olympus FV3000, DSS Imagetech, Bangalore) under a 10x, 20x and 40x DIC objective. Using the 3d reconstruction of *R. pomonella* brain as a guide (3) different brain regions were visualized and identified in FIJI open-source software (4).

**Quantification of neurotransmitters from the R. *pomonella* brain**

The level of neuromodulator per sample was calculated by a ratio of mass area of unknown sample and internal standards vs. ratio of mass area of internal standards and standards of each compound. The amount of each chemical (in ng) in the unknown sample was calculated from calibration curves. Further, outliers were identified and removed by calculating inter-quartile range (difference between Q1 and Q3 quartile) in Microsoft Excel for each age group. The lower bound was calculated by multiplying the IQR value by 1.5 then subtracting it from the Q1 data point. The upper bound was calculated by multiplying the IQR by 1.5 and adding it to the Q3 data point. In brief, the values lying 1.5 times below or above the IQR were considered outliers and removed from the further analysis. Quantities of each compound in each sample group were analysed using linear mixed model effects as mentioned above. In the model, age and host race were used as a fixed effect, whereas mass spec batch was used as a random effect. The satterthwaite approximations in the lmerTest package was used to test the significance of the effect. Further Tukey's HSD tests were performed for pairwise multiple comparisons with significance cut offs of 0.0001,0.001, and 0.05.

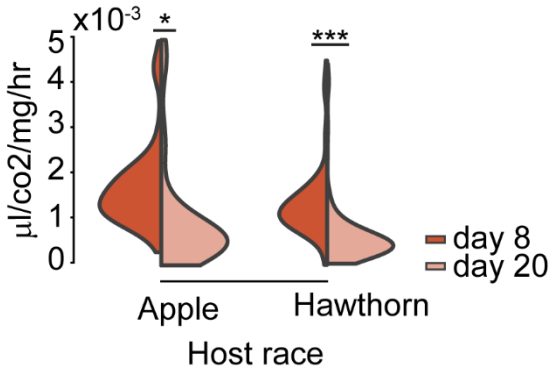



**Figure s1: Violin plots showing metabolic rates of pupae at two-time points for both host races**. Apple race, day 8, n=142, diapausing pupae; n=68, non-diapausing pupae; day 20, n= 145, diapausing pupae, n=62, non-diapausing pupae; hawthorn race, day 8, n=145, diapausing pupae; n=59, non-diapausing pupae; day 20, n= 152, diapausing pupae, n=17, non-diapausing pupae. Analysis performed using a linear mixed model to test for a difference in metabolic rates between day 8 and day 20. P value is significant < 0.05*, < 0.01**, < 0.001***

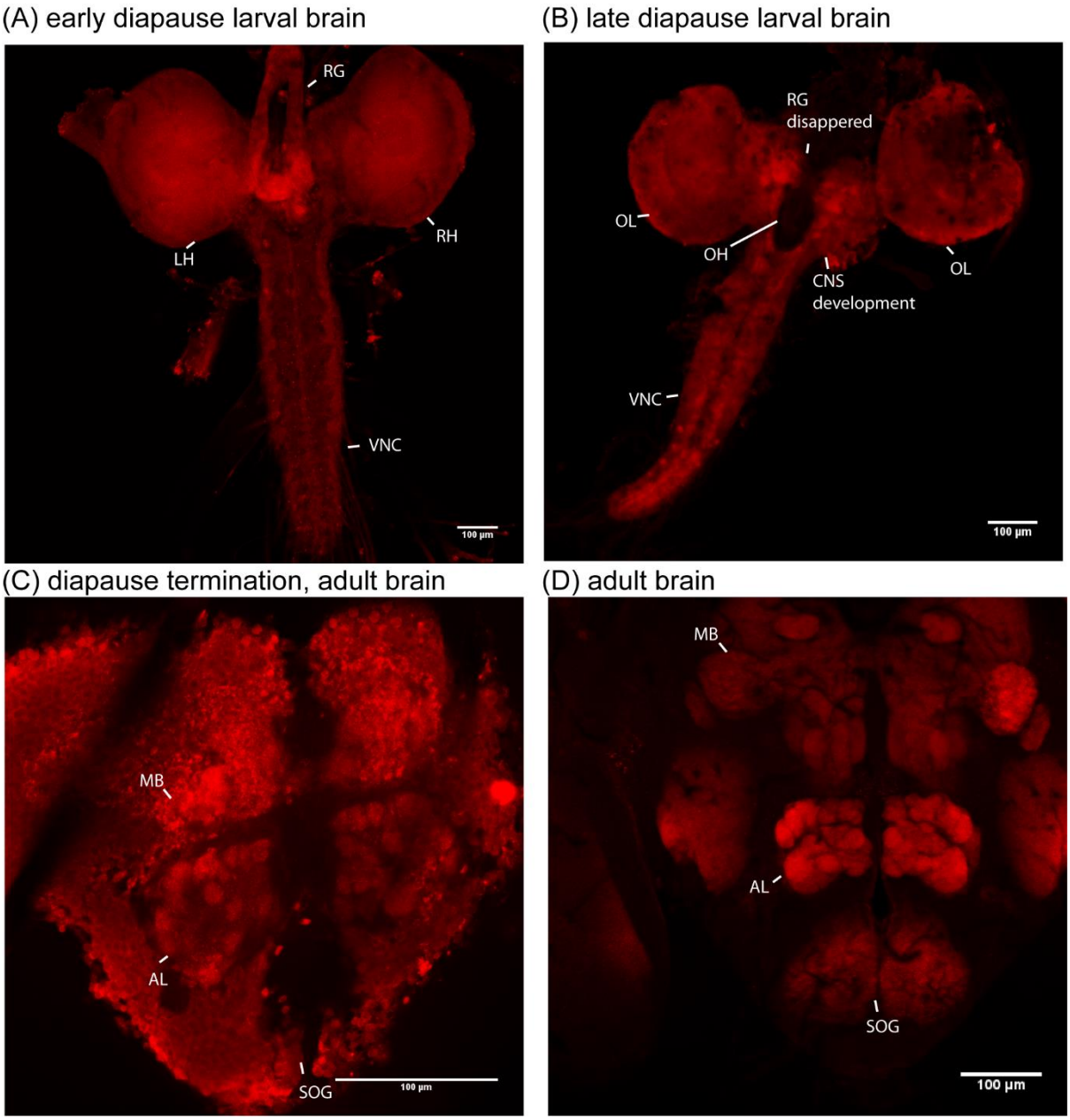



**Figure s2: Pupal brain morphology across developmental stages.** Confocal micrographs (1 µm depth) showing nc82 (red) staining of pupal brains at different stages. These are the same confocal images represented in Figure 1. The four images provide (A) an overview of the larval brain at diapause initiation (stage1, Figure 1) with two, left and right brain hemispheres (LH and RH), the ring gland (RG) and ventral nerve code (VNC), (B) the intermediate developing pupal brain at late diapause (stage 3) with clear formation of the optic lobe (OL), disappearance of the ring gland, oesophageal hole (OH) with CNS development, and VNC, (C) early adult brain (stage 4) following diapause termination with the first appearance of the boundaries of brain region like the antennal lobes (AL), mushroom bodies (MB), and suboesophageal ganglion (SOG), and (D) the adult brain (stage 7) with completely developed CNS after adult eclosion.

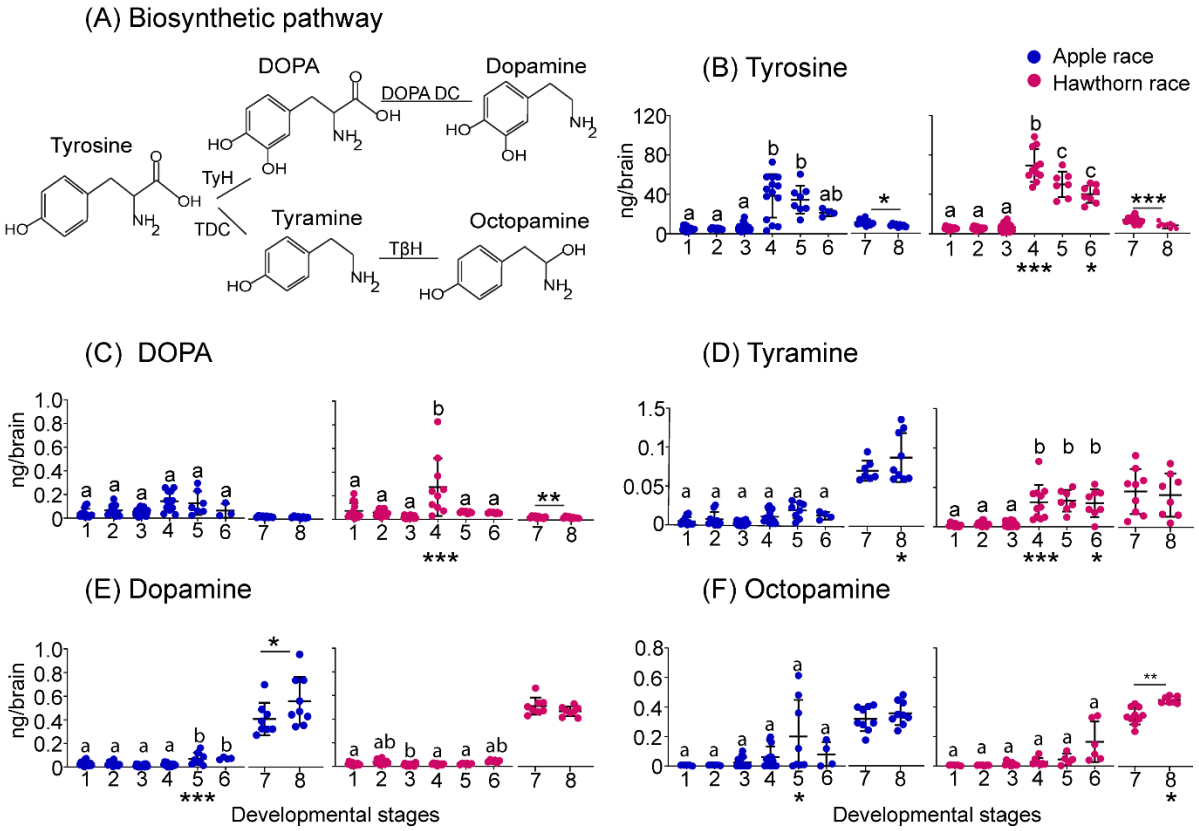



**Figure s3: Quantification of biogenic amines and their precursors across the different stages of development as defined in Figure 1.** (A) Biosynthetic pathways for octopamine and dopamine with enzymes, tyrosine hydroxylase (TyH), DOPA decarboxylase (DOPA DC), tyrosine decarboxylase and tyramine beta-hydroxylase (T□H); (B-F) Plots of neurochemical titres for both host races at different developmental stages with 4-15 samples per stage containing a pool of five brains in each sample, symbols represent medians with 95% CI: B) tyrosine; (C) DOPA; (D) tyramine; (E) dopamine; (F) octopamine. Letters below x-axis highlight specific developmental stages: N, onset of adult brain neurogenesis, D, development and differentiation of adult brain associated with entering the pharate-adult phase, and E, adult eclosion. Letters above each stage indicate similar titres of mentioned neurochemicals between pupal and pharate-adult stages within each host race ($P > 0,05$), and asterisks indicate differences in neurochemical titre between the two adult fly stages – pre-maturation and after sexual maturation. Asterisks below the x-axis indicate differences between host races at the equivalent stage of brain development. P-values represented are $< 0.05$ *, $< 0.01$ **, and $< 0.001$ ***, linear mixed effect model, followed by Tukey's HSD correction for multiple comparisons.

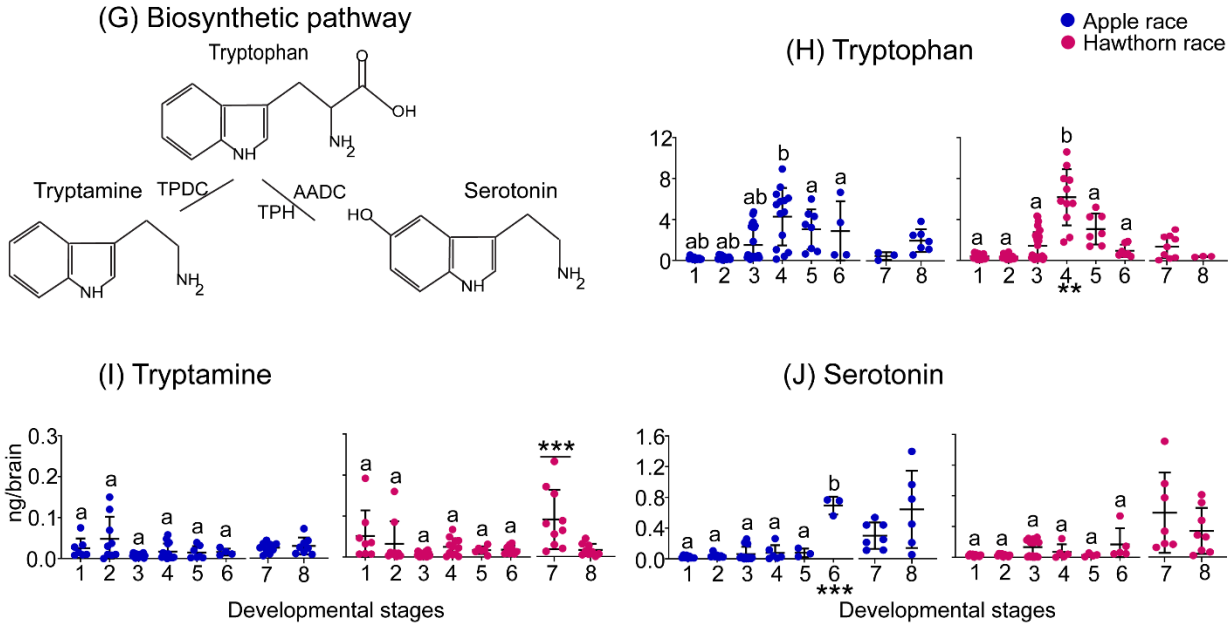



**Figure s4**: **Quantification of biogenic amines and their precursors across the different stages of development as defined in Figure 1.** (G) Biosynthetic pathway for serotonin with enzymes tryptophan decarboxylase (TPDC), tryptophan hydroxylase (TPH) and amino acid decarboxylase (AADC); (H-J) Scatter plots of neurochemical titres for both host races at different developmental stages with 4-15 samples per stage containing five brains each, symbols represent median with 95% CI: (H) tryptophan; (I) tryptamine; (J) serotonin. Lettering, asterisks, and statistical comparisons as in Figure s3.

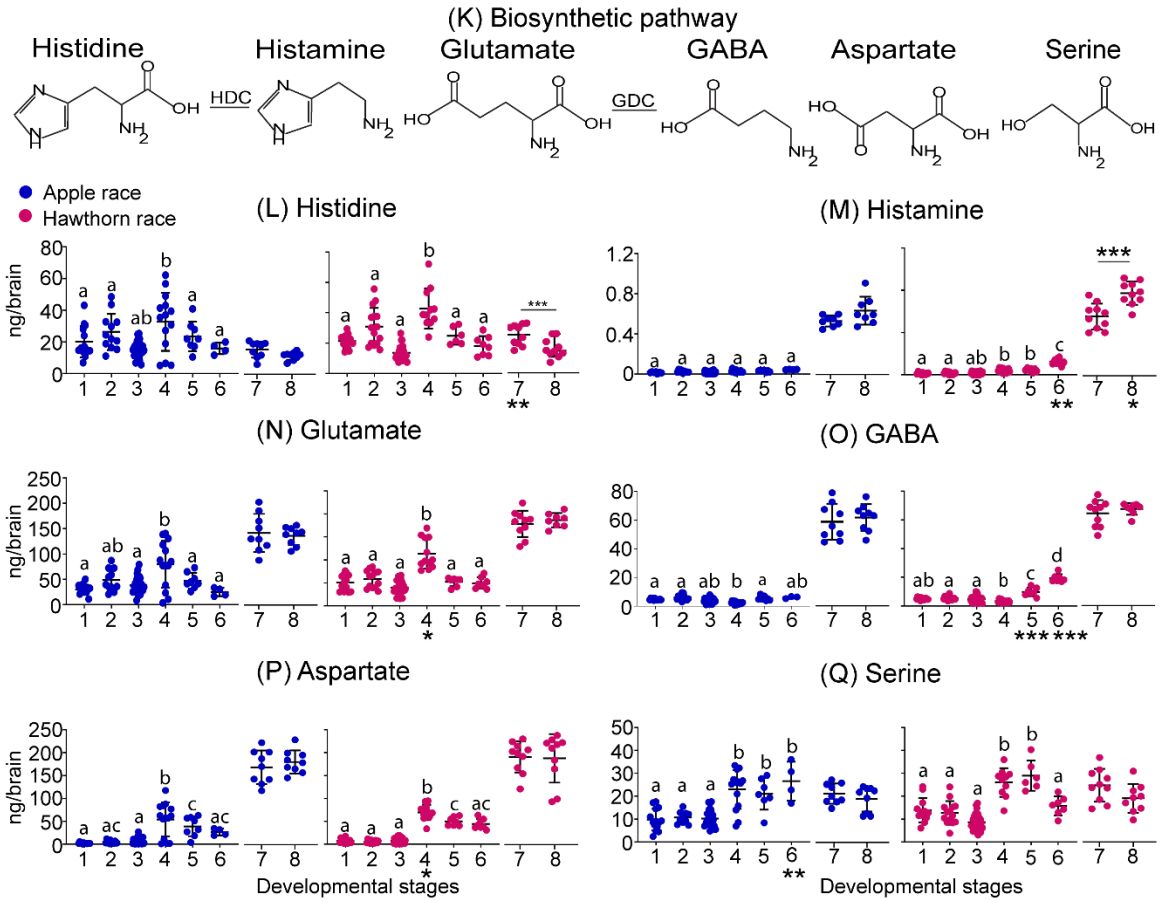
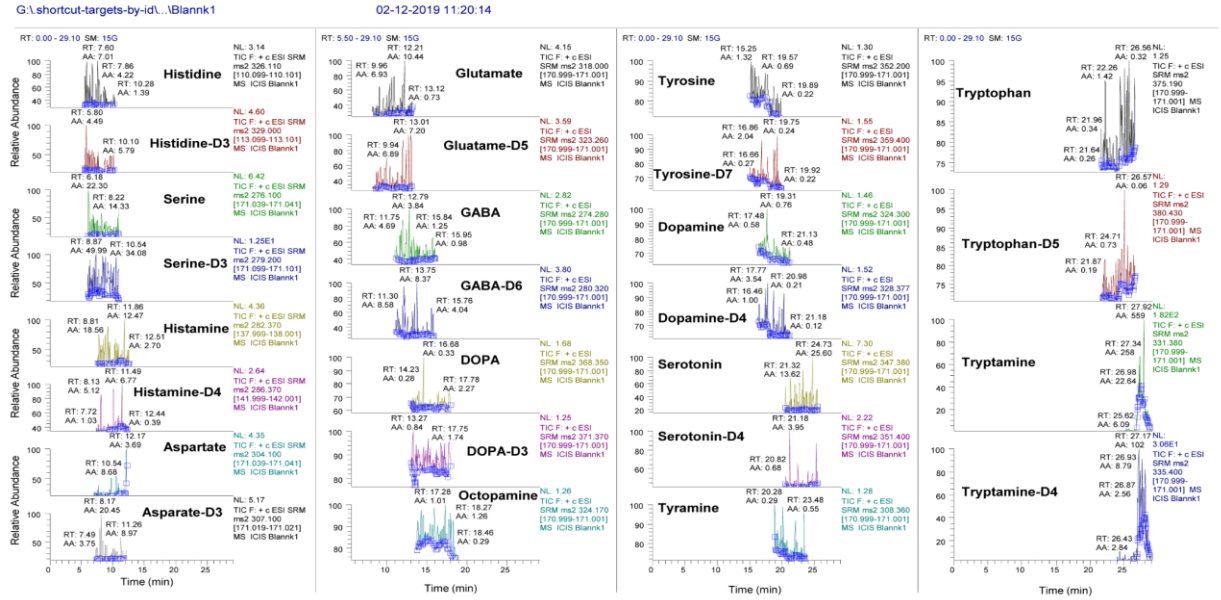
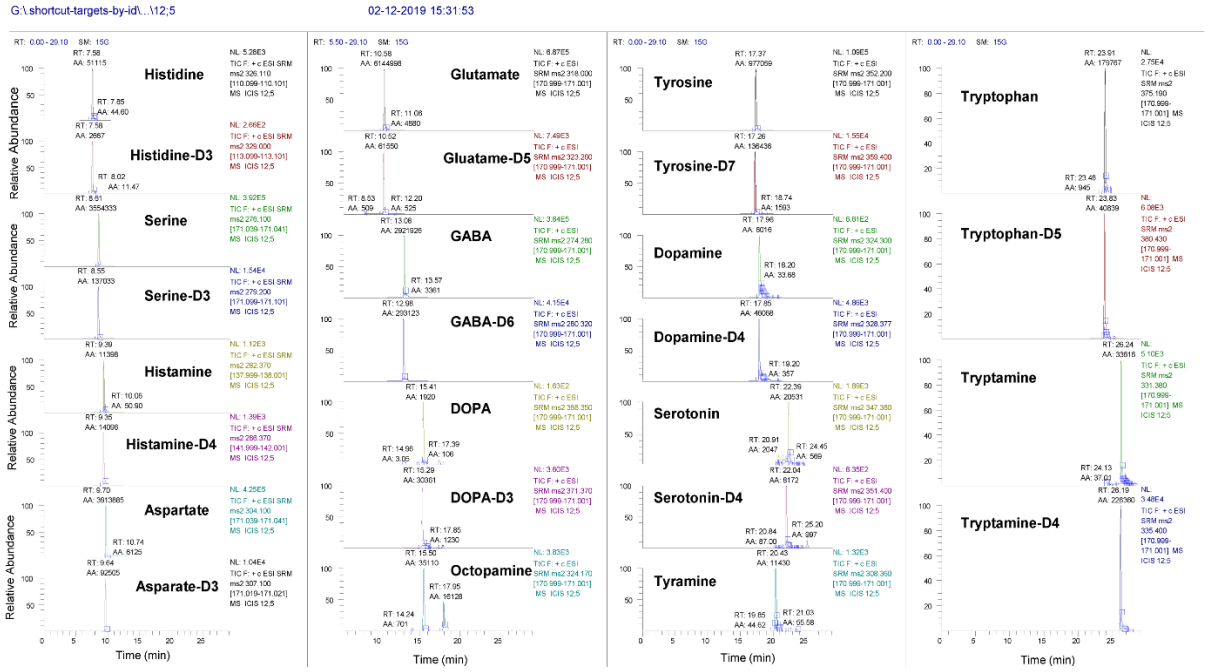



**Figure s5: Quantification of biogenic amines and their precursors across the different stages of development as defined in Figure 1.** (K) Biosynthetic pathway for histamine, GABA, aspartate and glutamate with enzymes histidine decarboxylase (HDC) and glutamic acid decarboxylase (GDC); (L-Q) Scatter plots of neurochemical titres for both host races at different developmental stages with 4-15 samples per stage containing five brains each, symbols represent median with 95% CI: (L) histidine; (M) histamine; (N) glutamate; (O) GABA; (P) aspartate; (Q) serine. Lettering, asterisks, and statistical comparisons as in Figure 3.

**A)**


G:\.shortcut-targets-by-id\...\Blannk1
02-12-2019 11:20:14
Histidine
Histidine-D3
Serine
Serine-D3
Histamine
Histamine-D4
Aspartate
Asparate-D3
Glutamate
Gluatame-D5
GABA
GABA-D6
DOPA
DOPA-D3
Octopamine
Tyrosine
Tyrosine-D7
Dopamine
Dopamine-D4
Serotonin
Serotonin-D4
Tyramine
Tryptophan
Tryptophan-D5
Tryptamine
Tryptamine-D4
Relative Abundance
Time (min)

**B)**


G:\.shortcut-targets-by-id\...\12;5
02-12-2019 15:31:53
Histidine
Histidine-D3
Serine
Serine-D3
Histamine
Histamine-D4
Aspartate
Asparate-D3
Glutamate
Gluatame-D5
GABA
GABA-D6
DOPA
DOPA-D3
Octopamine
Tyrosine
Tyrosine-D7
Dopamine
Dopamine-D4
Serotonin
Serotonin-D4
Tyramine
Tryptophan
Tryptophan-D5
Tryptamine
Tryptamine-D4
Relative Abundance
Time (min)

**c)**

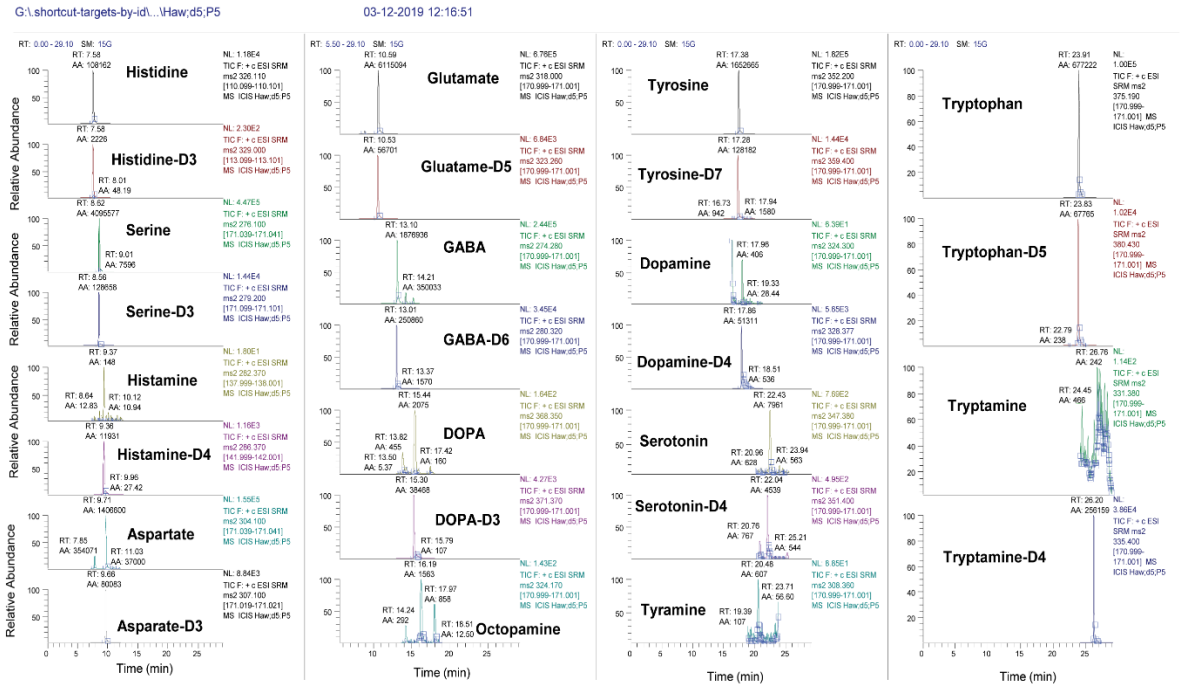



**Figure s6: Sample chromatograms obtained from LCMS neurochemical analysis.**

A) Blank (2% ACN prepared in 0.5% FA), B) Standard 12;5 from calibration curve and C) Pooled Hawthorn brain sample. For each neurochemical compound, the molecular weight and retention time details mention in table s11. Note that the blank chromatograms (A) do not show any peak at respected retention times. In contrast, calibration and sample chromatogram indicate the internal standard and compound peaks at particular retention time points where they are eluting.

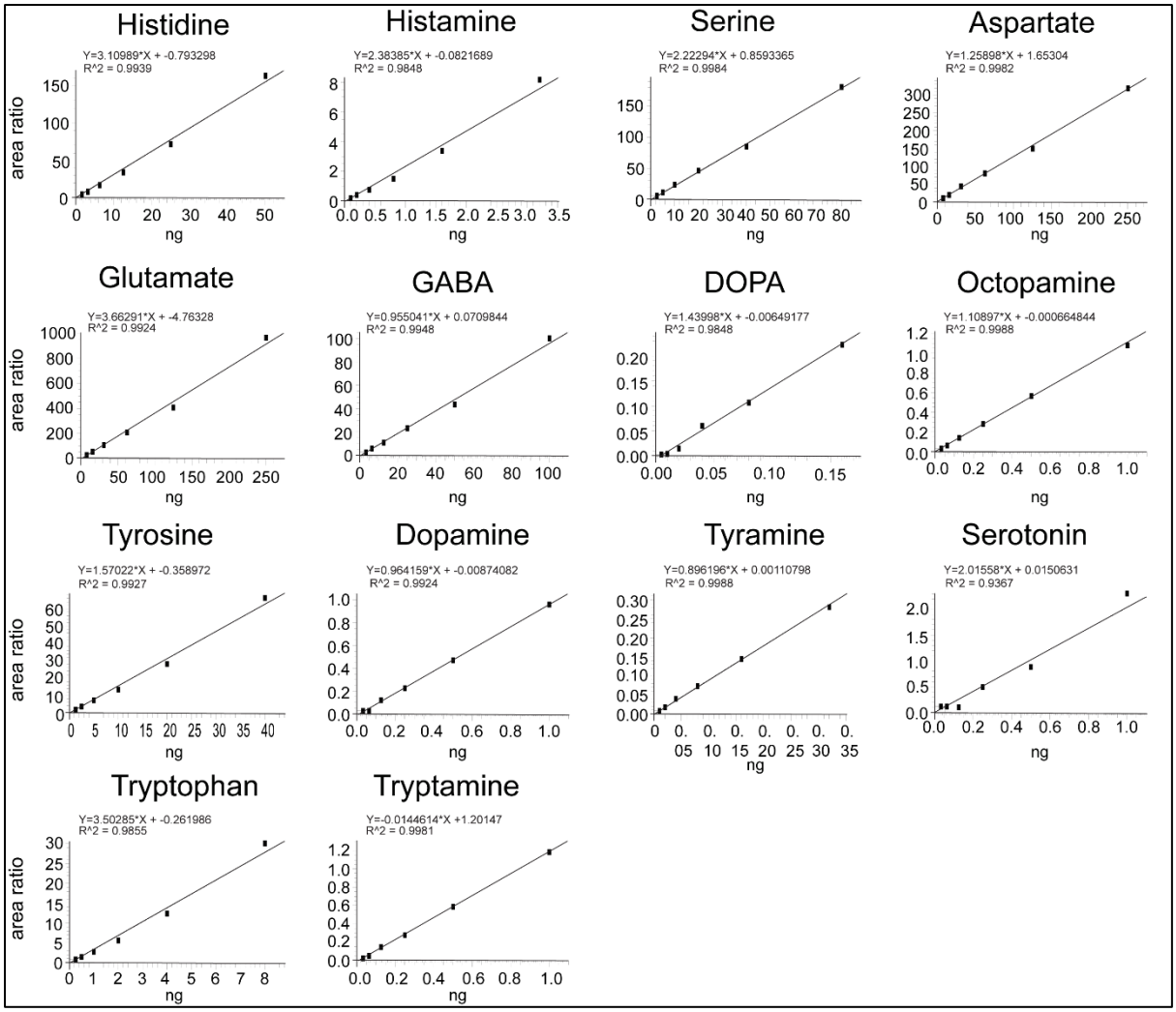



**Figure s7: Sample calibration curves for one of the LC-MS runs showing all 14 neurotransmitters analysed.**

**Table s1: Total number of pupae sampled after winter in the apple race**

| Total no. of | D 0 | D 5 | D 10 | D 15 | D 20 | D 25 | D 30 | D 35 | D 40 | D 45 | D 50 | D 55 | D 60 | D 65 | D 70 | total |
|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|
| pupae opened | 27 | 46 | 46 | 70 | 65 | 60 | 37 | 37 | 10 | 0 | 6 | 0 | 3 | 0 | 0 | 407 |
| larval brain,stage3 | 9 | 24 | 30 | 20 | 24 | 11 | 11 | 5 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 134 |
| adult brain, stage 4 | 11 | 1 | 1 | 3 | 6 | 20 | 13 | 11 | 4 | 0 | 0 | 0 | 0 | 0 | 0 | 70 |
| adult brain, stage 5 | 3 | 8 | 2 | 2 | 4 | 4 | 8 | 6 | 3 | 0 | 3 | 0 | 0 | 0 | 0 | 43 |
| adult brain, stage 6 | 0 | 4 | 5 | 6 | 1 | 1 | 1 | 1 | 0 | 0 | 0 | 0 | 1 | 0 | 0 | 20 |
| dead fly pupae | 3 | 5 | 8 | 38 | 29 | 21 | 0 | 10 | 3 | 0 | 0 | 0 | 0 | 0 | 0 | 117 |
| dead fly | 0 | 0 | 0 | 0 | 0 | 0 | 3 | 1 | 0 | 0 | 3 | 0 | 1 | 0 | 0 | 8 |
| wasp larvae | 0 | 2 | 0 | 1 | 1 | 3 | 0 | 2 | 0 | 0 | 0 | 0 | 1 | 0 | 0 | 10 |
| adult wasp | 1 | 2 | 0 | 0 | 0 | 0 | 1 | 1 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 5 |

**Table s2: Total number of pupae sampled after winter in the hawthorn race**

| Total no. of | D 0 | D 5 | D 10 | D 15 | D 20 | D 25 | D 30 | D 35 | D 40 | D 45 | D 50 | D 55 | D 60 | D 65 | D 70 | total |
|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|
| pupae opened | 26 | 28 | 56 | 10 | 9 | 50 | 41 | 30 | 55 | 78 | 74 | 31 | 51 | 47 | 19 | 605 |
| larval brain,stage3 | 22 | 25 | 25 | 2 | 6 | 20 | 20 | 19 | 14 | 14 | 2 | 1 | 0 | 1 | 0 | 171 |
| adult brain, stage 4 | 1 | 0 | 0 | 0 | 0 | 0 | 0 | 1 | 13 | 13 | 15 | 4 | 8 | 2 | 3 | 60 |
| adult brain, stage 5 | 0 | 0 | 1 | 0 | 0 | 0 | 0 | 0 | 0 | 3 | 3 | 1 | 16 | 8 | 3 | 35 |
| adult brain, stage 6 | 0 | 0 | 1 | 0 | 0 | 0 | 0 | 0 | 0 | 1 | 4 | 5 | 10 | 15 | 5 | 41 |
| dead fly pupae | 0 | 1 | 0 | 0 | 0 | 2 | 3 | 2 | 14 | 18 | 16 | 4 | 1 | 8 | 1 | 70 |
| dead fly | 0 | 0 | 3 | 0 | 0 | 6 | 2 | 1 | 3 | 9 | 4 | 1 | 5 | 3 | 4 | 41 |
| wasp larvae | 2 | 2 | 15 | 1 | 0 | 9 | 7 | 3 | 4 | 14 | 16 | 7 | 6 | 10 | 3 | 99 |
| adult wasp | 1 | 0 | 11 | 7 | 3 | 13 | 9 | 4 | 7 | 6 | 14 | 8 | 5 | 0 | 0 | 88 |

**Table s3-s6**: Generalised linear mixed models, Tukey's HSD tests was performed for pairwise multiple comparisons with a significance cut off of 0.0001***, 0.001**, or 0.05* as indicated.

**Table s3: Within and between host race metabolic rate comparisons.**

| compariosns | estimate | SE | df | t.ratio | p.value | |
|---|---|---|---|---|---|---|
| Apple day 8 - Apple day 20 | -0.0013 | 0.0004 | 237.8199 | -3.1127 | 0.0111 | * |
| Apple day 20 - Hawthorn day 20 | 0.0016 | 0.0005 | 136.0076 | 3.5224 | 0.0032 | ** |
| Apple day 8 - Hawthorn day 8 | 0.0002 | 0.0004 | 75.1650 | 0.5492 | 0.9465 | |
| Hawthorn day 8 - Hawthorn day 20 | -0.0027 | 0.0004 | 784.8679 | -6.2869 | 0.0001 | *** |

**Table s4: Between host race developmental stage comparisons for each neurotransmitter.**

| Histidine | estimate | SE | df | t.ratio | p.value | |
|---|---|---|---|---|---|---|
| Apple stage 1 - Hawthorn stage 1 | -1.3238 | 3.4071 | 151.0829 | -0.3885 | 1.0000 | |
| Apple stage 2 - Hawthorn stage 2 | -4.2289 | 3.3472 | 150.2634 | -1.2634 | 0.9823 | |
| Apple stage 3 - Hawthorn stage 3 | 2.7567 | 2.2972 | 152.9702 | 1.2000 | 0.9883 | |
| Apple stage 4 - Hawthorn stage 4 | -9.1510 | 3.5061 | 154.1119 | -2.6100 | 0.2835 | |
| Apple stage 5 - Hawthorn stage 5 | -1.3582 | 4.7046 | 153.6632 | -0.2887 | 1.0000 | |
| Apple stage 6 - Hawthorn stage 6 | -1.4942 | 5.4154 | 157.3576 | -0.2759 | 1.0000 | |
| Apple stage 7 - Hawthorn stage 7 | -7.1651 | 2.0019 | 34.5272 | -3.5791 | 0.0055 | ** |
| Apple stage 8 - Hawthorn stage 8 | -1.5290 | 2.0606 | 35.3436 | -0.7420 | 0.8794 | |
| Serine | estimate | SE | df | t.ratio | p.value | |
| Apple stage 1 - Hawthorn stage 1 | -4.3476 | 1.8536 | 138.3411 | -2.3454 | 0.4502 | |
| Apple stage 2 - Hawthorn stage 2 | -1.8663 | 1.9644 | 140.9179 | -0.9501 | 0.9984 | |
| Apple stage 3 - Hawthorn stage 3 | 2.0409 | 1.2809 | 139.9661 | 1.5934 | 0.9087 | |
| Apple stage 4 - Hawthorn stage 4 | -2.2410 | 1.9622 | 141.3912 | -1.1421 | 0.9921 | |
| Apple stage 5 - Hawthorn stage 5 | -7.3387 | 2.6361 | 141.0016 | -2.7839 | 0.1984 | |
| Apple stage 6 - Hawthorn stage 6 | 11.6866 | 3.0346 | 146.8927 | 3.8511 | 0.0093 | ** |
| Apple stage 7 - Hawthorn stage 7 | -2.2881 | 2.1287 | 33.5397 | -1.0749 | 0.7069 | |
| Apple stage 8 - Hawthorn stage 8 | 1.4459 | 2.2274 | 34.2703 | 0.6491 | 0.9151 | |
| Histamine | estimate | SE | df | t.ratio | p.value | |
| Apple stage 1 - Hawthorn stage 1 | 0.0024 | 0.0052 | 123.6331 | 0.4560 | 1.0000 | |
| Apple stage 2 - Hawthorn stage 2 | 0.0075 | 0.0048 | 122.3740 | 1.5637 | 0.9186 | |
| Apple stage 3 - Hawthorn stage 3 | 0.0009 | 0.0032 | 125.6815 | 0.2744 | 1.0000 | |
| Apple stage 4 - Hawthorn stage 4 | -0.0107 | 0.0043 | 123.2558 | -2.4914 | 0.3553 | |
| Apple stage 5 - Hawthorn stage 5 | -0.0087 | 0.0058 | 124.1761 | -1.4890 | 0.9409 | |
| Apple stage 6 - Hawthorn stage 6 | -0.0775 | 0.0067 | 124.0605 | -11.5464 | 0.0001 | *** |

| | | | | | | |
|---|---|---|---|---|---|---|
| Apple stage 7 - Hawthorn stage 7 | -0.0231 | 0.0476 | 33.1454 | -0.4861 | 0.9617 | |
| Apple stage 8 - Hawthorn stage 8 | -0.1543 | 0.0502 | 35.5456 | -3.0755 | 0.0201 | * |
| | | | | | | |
| Aspartate | estimate | SE | df | t.ratio | p.value | |
| Apple stage 1 - Hawthorn stage 1 | -3.6951 | 3.9919 | 150.1434 | -0.9256 | 0.9988 | |
| Apple stage 2 - Hawthorn stage 2 | -0.8473 | 3.7653 | 148.7253 | -0.2250 | 1.0000 | |
| Apple stage 3 - Hawthorn stage 3 | 2.8912 | 2.6426 | 149.8157 | 1.0941 | 0.9945 | |
| Apple stage 4 - Hawthorn stage 4 | -14.2007 | 3.9688 | 150.4097 | -3.5781 | 0.0228 | * |
| Apple stage 5 - Hawthorn stage 5 | -4.4480 | 5.2702 | 149.2740 | -0.8440 | 0.9995 | |
| Apple stage 6 - Hawthorn stage 6 | -11.8885 | 6.1894 | 150.9730 | -1.9208 | 0.7444 | |
| Apple stage 7 - Hawthorn stage 7 | -8.6402 | 12.0732 | 34.0117 | -0.7156 | 0.8902 | |
| Apple stage 8 - Hawthorn stage 8 | 9.1575 | 12.4722 | 34.4754 | 0.7342 | 0.8826 | |
| | | | | | | |
| Glutamate | estimate | SE | df | t.ratio | p.value | |
| Apple stage 1 - Hawthorn stage 1 | -15.3468 | 7.3757 | 150.9329 | -2.0807 | 0.6371 | |
| Apple stage 2 - Hawthorn stage 2 | -11.1878 | 7.5039 | 150.0898 | -1.4909 | 0.9410 | |
| Apple stage 3 - Hawthorn stage 3 | 2.5626 | 4.8625 | 150.9665 | 0.5270 | 1.0000 | |
| Apple stage 4 - Hawthorn stage 4 | -30.9558 | 7.2703 | 152.3285 | -4.2578 | 0.0021 | ** |
| Apple stage 5 - Hawthorn stage 5 | -5.3290 | 9.5928 | 151.8868 | -0.5555 | 1.0000 | |
| Apple stage 6 - Hawthorn stage 6 | -24.0413 | 11.5010 | 153.2042 | -2.0904 | 0.6303 | |
| Apple stage 7 - Hawthorn stage 7 | -15.4941 | 10.3199 | 33.0780 | -1.5014 | 0.4481 | |
| Apple stage 8 - Hawthorn stage 8 | -21.4532 | 11.4093 | 34.6585 | -1.8803 | 0.2550 | |
| | | | | | | |
| GABA | estimate | SE | df | t.ratio | p.value | |
| Apple stage 1 - Hawthorn stage 1 | -0.3228 | 0.6631 | 149.8323 | -0.4868 | 1.0000 | |
| Apple stage 2 - Hawthorn stage 2 | 0.5101 | 0.6595 | 146.8546 | 0.7736 | 0.9998 | |
| Apple stage 3 - Hawthorn stage 3 | -0.8280 | 0.4311 | 147.5445 | -1.9207 | 0.7444 | |
| Apple stage 4 - Hawthorn stage 4 | -0.6508 | 0.6661 | 153.4659 | -0.9770 | 0.9980 | |
| Apple stage 5 - Hawthorn stage 5 | -4.3573 | 0.8868 | 150.7387 | -4.9138 | 0.0001 | *** |
| Apple stage 6 - Hawthorn stage 6 | -13.4738 | 1.1482 | 151.4194 | -11.7348 | 0.0000 | *** |
| Apple stage 7 - Hawthorn stage 7 | -3.4803 | 3.1175 | 32.8182 | -1.1164 | 0.6822 | |
| Apple stage 8 - Hawthorn stage 8 | -0.4529 | 3.4660 | 34.2058 | -0.1307 | 0.9992 | |
| | | | | | | |
| Dopa | estimate | SE | df | t.ratio | p.value | |
| Apple stage 1 - Hawthorn stage 1 | -0.0312 | 0.0264 | 149.9665 | -1.1818 | 0.9896 | |
| Apple stage 2 - Hawthorn stage 2 | 0.0066 | 0.0273 | 151.0489 | 0.2408 | 1.0000 | |
| Apple stage 3 - Hawthorn stage 3 | 0.0361 | 0.0180 | 150.3388 | 2.0058 | 0.6888 | |
| Apple stage 4 - Hawthorn stage 4 | -0.1378 | 0.0295 | 151.2967 | -4.6798 | 0.0004 | *** |
| Apple stage 5 - Hawthorn stage 5 | 0.0591 | 0.0368 | 151.8817 | 1.6048 | 0.9048 | |
| Apple stage 6 - Hawthorn stage 6 | 0.0076 | 0.0477 | 153.2333 | 0.1594 | 1.0000 | |

| | | | | | | |
|---|---|---|---|---|---|---|
| Apple stage 7 - Hawthorn stage 7 | -0.0009 | 0.0017 | 32.4581 | -0.5609 | 0.9429 | |
| Apple stage 8 - Hawthorn stage 8 | 0.0014 | 0.0019 | 33.3627 | 0.7705 | 0.8672 | |
| | | | | | | |
| Octopamine | estimate | SE | df | t.ratio | p.value | |
| Apple stage 1 - Hawthorn stage 1 | -0.0006 | 0.0476 | 76.5057 | -0.0126 | 1.0000 | |
| Apple stage 2 - Hawthorn stage 2 | 0.0000 | 0.0447 | 78.2365 | 0.0003 | 1.0000 | |
| Apple stage 3 - Hawthorn stage 3 | 0.0255 | 0.0349 | 83.5626 | 0.7329 | 0.9999 | |
| Apple stage 4 - Hawthorn stage 4 | 0.0562 | 0.0333 | 80.0402 | 1.6869 | 0.8688 | |
| Apple stage 5 - Hawthorn stage 5 | 0.1658 | 0.0452 | 83.0806 | 3.6674 | 0.0206 | * |
| Apple stage 6 - Hawthorn stage 6 | -0.0460 | 0.0511 | 81.5941 | -0.8994 | 0.9990 | |
| Apple stage 7 - Hawthorn stage 7 | -0.0331 | 0.0272 | 32.5550 | -1.2151 | 0.6219 | |
| Apple stage 8 - Hawthorn stage 8 | -0.1006 | 0.0304 | 34.5325 | -3.3061 | 0.0113 | * |
| | | | | | | |
| Tyrosine | estimate | SE | df | t.ratio | p.value | |
| Apple stage 1 - Hawthorn stage 1 | -0.8273 | 3.1443 | 143.1745 | -0.2631 | 1.0000 | |
| Apple stage 2 - Hawthorn stage 2 | -0.4517 | 3.5577 | 144.4147 | -0.1270 | 1.0000 | |
| Apple stage 3 - Hawthorn stage 3 | 2.0068 | 2.2841 | 150.3338 | 0.8786 | 0.9992 | |
| Apple stage 4 - Hawthorn stage 4 | -29.1204 | 3.3965 | 151.2312 | -8.5737 | 0.0001 | *** |
| Apple stage 5 - Hawthorn stage 5 | -13.2214 | 4.4727 | 145.9859 | -2.9560 | 0.1329 | |
| Apple stage 6 - Hawthorn stage 6 | -18.4293 | 5.2246 | 152.3163 | -3.5274 | 0.0267 | |
| Apple stage 7 - Hawthorn stage 7 | -1.3088 | 1.0043 | 33.5635 | -1.3032 | 0.5674 | |
| Apple stage 8 - Hawthorn stage 8 | 0.6938 | 1.0493 | 34.4397 | 0.6612 | 0.9109 | |
| | | | | | | |
| Dopamine | estimate | SE | df | t.ratio | p.value | |
| Apple stage 1 - Hawthorn stage 1 | 0.0095 | 0.0063 | 128.4363 | 1.5266 | 0.9304 | |
| Apple stage 2 - Hawthorn stage 2 | -0.0112 | 0.0067 | 131.8681 | -1.6715 | 0.8778 | |
| Apple stage 3 - Hawthorn stage 3 | 0.0030 | 0.0053 | 138.6630 | 0.5725 | 1.0000 | |
| Apple stage 4 - Hawthorn stage 4 | 0.0022 | 0.0069 | 131.5225 | 0.3136 | 1.0000 | |
| Apple stage 5 - Hawthorn stage 5 | 0.0440 | 0.0089 | 134.1271 | 4.9272 | 0.0001 | *** |
| Apple stage 6 - Hawthorn stage 6 | 0.0211 | 0.0108 | 135.5962 | 1.9635 | 0.7168 | |
| Apple stage 7 - Hawthorn stage 7 | -0.0829 | 0.0529 | 31.2879 | -1.5692 | 0.4102 | |
| Apple stage 8 - Hawthorn stage 8 | 0.0934 | 0.0512 | 31.1948 | 1.8223 | 0.2823 | |
| | | | | | | |
| Tyramine | estimate | SE | df | t.ratio | p.value | |
| Apple stage 1 - Hawthorn stage 1 | 0.0018 | 0.0031 | 137.0627 | 0.5680 | 1.0000 | |
| Apple stage 2 - Hawthorn stage 2 | 0.0045 | 0.0029 | 134.6750 | 1.5361 | 0.9277 | |
| Apple stage 3 - Hawthorn stage 3 | 0.0009 | 0.0020 | 135.1031 | 0.4848 | 1.0000 | |
| Apple stage 4 - Hawthorn stage 4 | -0.0173 | 0.0028 | 136.4020 | -6.1011 | 0.0001 | *** |
| Apple stage 5 - Hawthorn stage 5 | -0.0068 | 0.0037 | 136.8321 | -1.8537 | 0.7850 | |
| Apple stage 6 - Hawthorn stage 6 | -0.0154 | 0.0045 | 137.4406 | -3.4070 | 0.0393 | * |

| | | | | | | |
|---|---|---|---|---|---|---|
| Apple stage 7 - Hawthorn stage 7 | 0.0127 | 0.0110 | 31.2861 | 1.1570 | 0.6577 | |
| Apple stage 8 - Hawthorn stage 8 | 0.0358 | 0.0108 | 32.0899 | 3.3242 | 0.0114 | * |

| Serotonine | estimate | SE | df | t.ratio | p.value | |
|---|---|---|---|---|---|---|
| Apple stage 1 - Hawthorn stage 1 | 0.0009 | 0.0279 | 89.5853 | 0.0329 | 1.0000 | |
| Apple stage 2 - Hawthorn stage 2 | 0.0085 | 0.0291 | 85.6962 | 0.2903 | 1.0000 | |
| Apple stage 3 - Hawthorn stage 3 | -0.0133 | 0.0217 | 87.6624 | -0.6126 | 1.0000 | |
| Apple stage 4 - Hawthorn stage 4 | 0.0466 | 0.0416 | 94.9574 | 1.1204 | 0.9930 | |
| Apple stage 5 - Hawthorn stage 5 | 0.0660 | 0.0458 | 88.5421 | 1.4405 | 0.9519 | |
| Apple stage 6 - Hawthorn stage 6 | 0.4526 | 0.0462 | 91.0937 | 9.7915 | 0.0001 | *** |
| Apple stage 7 - Hawthorn stage 7 | -0.1029 | 0.1705 | 24.5415 | -0.6036 | 0.9299 | |
| Apple stage 8 - Hawthorn stage 8 | 0.3304 | 0.1702 | 26.8478 | 1.9415 | 0.2352 | |

| Tryptophan | estimate | SE | df | t.ratio | p.value | |
|---|---|---|---|---|---|---|
| Apple stage 1 - Hawthorn stage 1 | -0.1678 | 0.3689 | 153.5635 | -0.4549 | 1.0000 | |
| Apple stage 2 - Hawthorn stage 2 | -0.0997 | 0.3661 | 153.1281 | -0.2724 | 1.0000 | |
| Apple stage 3 - Hawthorn stage 3 | 0.4235 | 0.2410 | 153.2847 | 1.7576 | 0.8378 | |
| Apple stage 4 - Hawthorn stage 4 | -1.5851 | 0.3798 | 153.9048 | -4.1734 | 0.0028 | ** |
| Apple stage 5 - Hawthorn stage 5 | 0.0957 | 0.4913 | 153.9227 | 0.1947 | 1.0000 | |
| Apple stage 6 - Hawthorn stage 6 | 1.1234 | 0.6015 | 154.2040 | 1.8678 | 0.7770 | |
| Apple stage 7 - Hawthorn stage 7 | -1.0653 | 0.6399 | 22.0000 | -1.6650 | 0.3651 | |
| Apple stage 8 - Hawthorn stage 8 | 1.5244 | 0.6623 | 22.0000 | 2.3016 | 0.1283 | |

| Tryptamine | estimate | SE | df | t.ratio | p.value | |
|---|---|---|---|---|---|---|
| Apple stage 1 - Hawthorn stage 1 | -0.0155 | 0.0128 | 117.2359 | -1.2105 | 0.9872 | |
| Apple stage 2 - Hawthorn stage 2 | 0.0075 | 0.0116 | 116.3439 | 0.6443 | 1.0000 | |
| Apple stage 3 - Hawthorn stage 3 | 0.0004 | 0.0078 | 115.6457 | 0.0478 | 1.0000 | |
| Apple stage 4 - Hawthorn stage 4 | -0.0045 | 0.0097 | 117.6350 | -0.4698 | 1.0000 | |
| Apple stage 5 - Hawthorn stage 5 | 0.0040 | 0.0136 | 115.5865 | 0.2917 | 1.0000 | |
| Apple stage 6 - Hawthorn stage 6 | 0.0075 | 0.0154 | 118.8689 | 0.4883 | 1.0000 | |
| Apple stage 7 - Hawthorn stage 7 | -0.0566 | 0.0167 | 33.8268 | -3.3831 | 0.0094 | ** |
| Apple stage 8 - Hawthorn stage 8 | 0.0235 | 0.0183 | 35.9997 | 1.2844 | 0.5786 | |

**Table s5: Apple race developmental stage comparisons for each neurotransmitter.**

| Histidine | estimate | SE | df | t.ratio | p.value | |
|---|---|---|---|---|---|---|
| Stage 1 - stage 2 | 6.3710 | 3.4071 | 151.0829 | 1.8699 | 0.7757 | |
| Stage 1 - stage 3 | -4.0614 | 3.5846 | 33.8738 | -1.1330 | 0.9907 | |
| Stage 1 - stage 4 | -10.7407 | 3.8992 | 47.0710 | -2.7546 | 0.2328 | |
| Stage 1 - stage 5 | 1.3674 | 4.3923 | 66.0851 | 0.3113 | 1.0000 | |
| Stage 1 - stage 6 | -4.7250 | 5.4277 | 104.0757 | -0.8705 | 0.9993 | |
| Stage 2 - stage 3 | 10.4324 | 3.7043 | 38.2197 | 2.8163 | 0.2135 | |
| Stage 2 - stage 4 | -4.3698 | 4.0095 | 51.7618 | -1.0898 | 0.9939 | |
| Stage 2 - stage 5 | 5.0036 | 4.4905 | 70.5918 | 1.1143 | 0.9930 | |
| Stage 2 - stage 6 | 11.0960 | 5.5075 | 107.2170 | 2.0147 | 0.6825 | |
| Stage 3 - stage 4 | -14.8022 | 3.0304 | 161.9721 | -4.8845 | 0.0002 | *** |
| Stage 3 - stage 5 | 5.4288 | 3.6700 | 161.9028 | 1.4792 | 0.9442 | |
| Stage 3 - stage 6 | -0.6636 | 4.8474 | 161.6184 | -0.1369 | 1.0000 | |
| Stage 4 - stage 5 | -9.3734 | 3.8692 | 154.6094 | -2.4226 | 0.3975 | |
| Stage 4 - stage 6 | -15.4657 | 5.0646 | 158.8954 | -3.0537 | 0.1032 | |
| Stage 5 - stage 6 | -6.0924 | 5.3788 | 154.4870 | -1.1327 | 0.9927 | |
| Stage 6 - stage 7 | 1.2020 | 5.4286 | 126.1111 | 0.2214 | 1.0000 | |
| Stage 7 - stage 8 | -3.8320 | 2.0147 | 34.3189 | -1.9020 | 0.2461 | |
| **Serine** | **estimate** | **SE** | **df** | **t.ratio** | **p.value** | |
| Stage 1 - stage 2 | 1.0629 | 1.9492 | 138.0375 | 0.5453 | 1.0000 | |
| Stage 1 - stage 3 | 1.0320 | 2.0756 | 22.0955 | 0.4972 | 1.0000 | |
| Stage 1 - stage 4 | -12.1406 | 2.2253 | 30.0121 | -5.4557 | 0.0003 | *** |
| Stage 1 - stage 5 | 10.0534 | 2.5654 | 47.8072 | 3.9189 | 0.0134 | * |
| Stage 1 - stage 6 | 16.2475 | 3.0339 | 76.3100 | 5.3553 | 0.0001 | *** |
| Stage 2 - stage 3 | 0.0309 | 2.2305 | 28.7229 | 0.0139 | 1.0000 | |
| Stage 2 - stage 4 | -11.0777 | 2.3705 | 37.3524 | -4.6732 | 0.0020 | ** |
| Stage 2 - stage 5 | -8.9904 | 2.6922 | 55.3685 | -3.3394 | 0.0602 | |
| Stage 2 - stage 6 | -15.1846 | 3.1419 | 82.7951 | -4.8329 | 0.0004 | *** |
| Stage 3 - stage 4 | -11.1086 | 1.6765 | 152.9994 | -6.6259 | 0.0001 | *** |
| Stage 3 - stage 5 | 9.0214 | 2.1364 | 152.8918 | 4.2227 | 0.0024 | ** |
| Stage 3 - stage 6 | 15.2155 | 2.6621 | 151.7478 | 5.7157 | 0.0001 | *** |

| | | | | | | |
|---|---|---|---|---|---|---|
| Stage 4 - stage 5 | -2.0872 | 2.2173 | 144.3248 | -0.9413 | 0.9985 | |
| Stage 4 - stage 6 | 4.1069 | 2.7698 | 147.4602 | 1.4828 | 0.9431 | |
| Stage 5 - stage 6 | 6.1941 | 2.9927 | 141.4463 | 2.0697 | 0.6448 | |
| Stage 6 - stage 7 | -5.0133 | 3.3908 | 83.9981 | -1.4785 | 0.9821 | |
| Stage 7 - stage 8 | -1.9049 | 2.1402 | 33.2748 | -0.8900 | 0.8100 | |
| Histamine | estimate | SE | df | t.ratio | p.value | |
| Stage 1 - stage 2 | 0.0081 | 0.0054 | 124.0529 | 1.4793 | 0.9435 | |
| Stage 1 - stage 3 | 0.0044 | 0.0073 | 22.0036 | 0.5997 | 1.0000 | |
| Stage 1 - stage 4 | -0.0114 | 0.0076 | 25.0608 | -1.5019 | 0.9265 | |
| Stage 1 - stage 5 | 0.0120 | 0.0080 | 31.2531 | 1.4888 | 0.9328 | |
| Stage 1 - stage 6 | 0.0187 | 0.0088 | 43.3008 | 2.1158 | 0.6149 | |
| Stage 2 - stage 3 | 0.0037 | 0.0069 | 18.5033 | 0.5353 | 1.0000 | |
| Stage 2 - stage 4 | -0.0033 | 0.0072 | 21.4419 | -0.4550 | 1.0000 | |
| Stage 2 - stage 5 | -0.0039 | 0.0077 | 27.4179 | -0.5041 | 1.0000 | |
| Stage 2 - stage 6 | -0.0107 | 0.0086 | 39.3271 | -1.2439 | 0.9814 | |
| Stage 3 - stage 4 | -0.0070 | 0.0040 | 128.5870 | -1.7647 | 0.8337 | |
| Stage 3 - stage 5 | 0.0076 | 0.0048 | 126.9465 | 1.5858 | 0.9111 | |
| Stage 3 - stage 6 | 0.0144 | 0.0061 | 126.4946 | 2.3716 | 0.4329 | |
| Stage 4 - stage 5 | 0.0006 | 0.0050 | 123.0837 | 0.1194 | 1.0000 | |
| Stage 4 - stage 6 | 0.0074 | 0.0063 | 124.2264 | 1.1637 | 0.9907 | |
| Stage 5 - stage 6 | 0.0068 | 0.0068 | 122.5118 | 1.0012 | 0.9974 | |
| Stage 6 - stage 7 | 0.5474 | 0.0367 | 65.1478 | 14.9044 | 0.0001 | *** |
| Stage 7 - stage 8 | 0.1078 | 0.0490 | 32.2913 | 2.1992 | 0.1450 | |
| Aspartate | estimate | SE | df | t.ratio | p.value | |
| Stage 1 - stage 2 | 1.3990 | 3.9919 | 150.1434 | 0.3505 | 1.0000 | |
| Stage 1 - stage 3 | 9.2458 | 7.3079 | 14.8824 | 1.2652 | 0.9720 | |
| Stage 1 - stage 4 | -40.6847 | 7.4747 | 16.3835 | -5.4430 | 0.0021 | ** |
| Stage 1 - stage 5 | 29.2739 | 7.9529 | 20.6859 | 3.6809 | 0.0481 | * |
| Stage 1 - stage 6 | 17.3012 | 8.6076 | 28.0209 | 2.0100 | 0.6836 | |
| Stage 2 - stage 3 | -7.8468 | 7.2160 | 14.2074 | -1.0874 | 0.9904 | |
| Stage 2 - stage 4 | -39.2857 | 7.3849 | 15.6796 | -5.3198 | 0.0029 | ** |

| | | | | | | |
|---|---|---|---|---|---|---|
| Stage 2 - stage 5 | -27.8749 | 7.8686 | 19.9217 | -3.5426 | 0.0655 | |
| Stage 2 - stage 6 | -15.9022 | 8.5297 | 27.1797 | -1.8643 | 0.7697 | |
| | | | | | | |
| Stage 3 - stage 4 | -31.4389 | 3.7533 | 156.0706 | -8.3763 | 0.0001 | *** |
| Stage 3 - stage 5 | 20.0281 | 4.7937 | 158.0994 | 4.1780 | 0.0028 | ** |
| Stage 3 - stage 6 | 8.0554 | 5.6950 | 154.3405 | 1.4145 | 0.9591 | |
| | | | | | | |
| Stage 4 - stage 5 | -11.4108 | 4.6537 | 150.4490 | -2.4520 | 0.3785 | |
| Stage 4 - stage 6 | -23.3835 | 5.8111 | 150.4609 | -4.0239 | 0.0050 | ** |
| | | | | | | |
| Stage 5 - stage 6 | -11.9727 | 6.2388 | 149.3851 | -1.9191 | 0.7455 | |
| Stage 6 - stage 7 | 140.8102 | 15.4588 | 28.1886 | 9.1088 | 0.0001 | *** |
| | | | | | | |
| Stage 7 - stage 8 | 19.7866 | 12.1420 | 33.9396 | 1.6296 | 0.3761 | |
| | | | | | | |
| Glutamate | estimate | SE | df | t.ratio | p.value | |
| Stage 1 - stage 2 | 14.4270 | 7.6977 | 151.1464 | 1.8742 | 0.7731 | |
| Stage 1 - stage 3 | 8.1177 | 9.8073 | 23.4441 | 0.8277 | 0.9993 | |
| Stage 1 - stage 4 | -38.4449 | 10.2692 | 28.3157 | -3.7437 | 0.0321 | * |
| Stage 1 - stage 5 | 5.2307 | 11.1385 | 37.5008 | 0.4696 | 1.0000 | |
| Stage 1 - stage 6 | -14.6877 | 13.0425 | 61.9126 | -1.1261 | 0.9922 | |
| | | | | | | |
| Stage 2 - stage 3 | 6.3093 | 9.9010 | 24.4487 | 0.6372 | 0.9999 | |
| Stage 2 - stage 4 | -24.0178 | 10.3587 | 29.4102 | -2.3186 | 0.4864 | |
| Stage 2 - stage 5 | 9.1963 | 11.2211 | 38.6990 | 0.8196 | 0.9995 | |
| Stage 2 - stage 6 | 29.1148 | 13.1131 | 63.2049 | 2.2203 | 0.5418 | |
| | | | | | | |
| Stage 3 - stage 4 | -30.3272 | 6.6180 | 159.0077 | -4.5825 | 0.0006 | *** |
| Stage 3 - stage 5 | -2.8870 | 7.9892 | 159.6720 | -0.3614 | 1.0000 | |
| Stage 3 - stage 6 | -22.8054 | 10.4302 | 157.5626 | -2.1865 | 0.5619 | |
| | | | | | | |
| Stage 4 - stage 5 | -33.2141 | 8.1954 | 151.8280 | -4.0528 | 0.0045 | ** |
| Stage 4 - stage 6 | -53.1326 | 10.7735 | 153.8021 | -4.9318 | 0.0001 | *** |
| | | | | | | |
| Stage 5 - stage 6 | -19.9185 | 11.3870 | 151.1367 | -1.7492 | 0.8420 | |
| Stage 6 - stage 7 | 138.7044 | 14.7222 | 70.0816 | 9.4215 | 0.0001 | *** |
| | | | | | | |
| Stage 7 - stage 8 | -5.3597 | 10.4487 | 33.5306 | -0.5130 | 0.9554 | |
| | | | | | | |
| GABA | estimate | SE | df | t.ratio | p.value | |
| Stage 1 - stage 2 | 1.2415 | 0.6770 | 150.3439 | 1.8337 | 0.7968 | |

| | | | | | | |
|---|---|---|---|---|---|---|
| Stage 1 - stage 3 | -0.6305 | 0.6577 | 48.2274 | -0.9586 | 0.9979 | |
| Stage 1 - stage 4 | 2.4515 | 0.7364 | 69.1423 | 3.3288 | 0.0577 | * |
| Stage 1 - stage 5 | 0.5087 | 0.8492 | 95.4747 | 0.5990 | 1.0000 | |
| Stage 1 - stage 6 | 0.9161 | 1.1295 | 130.6834 | 0.8111 | 0.9996 | |
| | | | | | | |
| Stage 2 - stage 3 | 1.8720 | 0.6565 | 48.5321 | 2.8514 | 0.1915 | |
| Stage 2 - stage 4 | 3.6929 | 0.7354 | 69.5930 | 5.0219 | 0.0002 | *** |
| Stage 2 - stage 5 | 0.7328 | 0.8482 | 96.0020 | 0.8639 | 0.9993 | |
| Stage 2 - stage 6 | 0.3254 | 1.1288 | 131.0409 | 0.2883 | 1.0000 | |
| | | | | | | |
| Stage 3 - stage 4 | 1.8209 | 0.5774 | 157.9992 | 3.1534 | 0.0793 | |
| Stage 3 - stage 5 | 1.1392 | 0.7232 | 157.5404 | 1.5752 | 0.9155 | |
| Stage 3 - stage 6 | 1.5466 | 1.0421 | 157.3267 | 1.4842 | 0.9429 | |
| | | | | | | |
| Stage 4 - stage 5 | 2.9601 | 0.7860 | 155.4472 | 3.7659 | 0.0122 | * |
| Stage 4 - stage 6 | 3.3675 | 1.0924 | 157.9889 | 3.0826 | 0.0958 | |
| | | | | | | |
| Stage 5 - stage 6 | 0.4074 | 1.1487 | 151.7796 | 0.3547 | 1.0000 | |
| Stage 6 - stage 7 | 54.6217 | 3.1419 | 48.2363 | 17.3849 | 0.0001 | *** |
| | | | | | | |
| Stage 7 - stage 8 | 3.1468 | 3.1622 | 33.4048 | 0.9951 | 0.7532 | |

| Dopa | estimate | SE | df | t.ratio | p.value |
|---|---|---|---|---|---|
| Stage 1 - stage 2 | 0.0226 | 0.0277 | 151.2025 | 0.8146 | 0.9996 |
| Stage 1 - stage 3 | 0.0100 | 0.0262 | 51.4381 | 0.3824 | 1.0000 |
| Stage 1 - stage 4 | -0.0903 | 0.0300 | 76.8582 | -3.0122 | 0.1250 |
| Stage 1 - stage 5 | 0.0839 | 0.0344 | 99.0306 | 2.4369 | 0.3920 |
| Stage 1 - stage 6 | 0.0381 | 0.0461 | 135.8433 | 0.8257 | 0.9996 |
| | | | | | |
| Stage 2 - stage 3 | 0.0126 | 0.0280 | 61.1593 | 0.4493 | 1.0000 |
| Stage 2 - stage 4 | -0.0677 | 0.0316 | 85.0699 | -2.1440 | 0.5935 |
| Stage 2 - stage 5 | -0.0613 | 0.0358 | 104.8080 | -1.7109 | 0.8592 |
| Stage 2 - stage 6 | -0.0155 | 0.0472 | 137.5124 | -0.3286 | 1.0000 |
| | | | | | |
| Stage 3 - stage 4 | -0.0803 | 0.0250 | 156.1366 | -3.2082 | 0.0684 |
| Stage 3 - stage 5 | 0.0739 | 0.0305 | 154.7410 | 2.4243 | 0.3964 |
| Stage 3 - stage 6 | 0.0281 | 0.0433 | 157.5659 | 0.6487 | 1.0000 |
| | | | | | |
| Stage 4 - stage 5 | -0.0064 | 0.0328 | 152.0740 | -0.1953 | 1.0000 |
| Stage 4 - stage 6 | -0.0522 | 0.0454 | 157.3919 | -1.1481 | 0.9919 |

| | | | | | | |
|---|---|---|---|---|---|---|
| Stage 5 - stage 6 | -0.0458 | 0.0478 | 153.4026 | -0.9575 | 0.9983 | |
| Stage 6 - stage 7 | -0.0595 | 0.0438 | 164.5738 | -1.3589 | 0.9928 | |
| Stage 7 - stage 8 | -0.0036 | 0.0017 | 32.3078 | -2.1342 | 0.1639 | |
| Octopamine | estimate | SE | df | t.ratio | p.value | |
| Stage 1 - stage 2 | 0.0012 | 0.0476 | 76.5057 | 0.0252 | 1.0000 | |
| Stage 1 - stage 3 | 0.0400 | 0.0579 | 14.2248 | 0.6908 | 0.9998 | |
| Stage 1 - stage 4 | -0.0636 | 0.0579 | 14.3025 | -1.0975 | 0.9897 | |
| Stage 1 - stage 5 | 0.1777 | 0.0600 | 16.7666 | 2.9586 | 0.2045 | |
| Stage 1 - stage 6 | 0.0462 | 0.0671 | 23.8007 | 0.6889 | 0.9999 | |
| Stage 2 - stage 3 | -0.0388 | 0.0579 | 14.2248 | -0.6700 | 0.9999 | |
| Stage 2 - stage 4 | -0.0624 | 0.0579 | 14.3025 | -1.0768 | 0.9911 | |
| Stage 2 - stage 5 | -0.1765 | 0.0600 | 16.7666 | -2.9386 | 0.2111 | |
| Stage 2 - stage 6 | -0.0450 | 0.0671 | 23.8007 | -0.6710 | 0.9999 | |
| Stage 3 - stage 4 | -0.0236 | 0.0303 | 79.8946 | -0.7775 | 0.9997 | |
| Stage 3 - stage 5 | 0.1376 | 0.0366 | 88.3850 | 3.7659 | 0.0148 | * |
| Stage 3 - stage 6 | 0.0062 | 0.0463 | 86.0316 | 0.1341 | 1.0000 | |
| Stage 4 - stage 5 | 0.1141 | 0.0355 | 85.9109 | 3.2161 | 0.0737 | |
| Stage 4 - stage 6 | -0.0174 | 0.0468 | 85.3827 | -0.3711 | 1.0000 | |
| Stage 5 - stage 6 | -0.1314 | 0.0482 | 82.6230 | -2.7294 | 0.2309 | |
| Stage 6 - stage 7 | 0.2658 | 0.0531 | 52.0130 | 5.0052 | 0.0007 | *** |
| Stage 7 - stage 8 | 0.0347 | 0.0275 | 31.0306 | 1.2607 | 0.5941 | |
| Tyrosine | estimate | SE | df | t.ratio | p.value | |
| Stage 1 - stage 2 | 0.2142 | 3.5059 | 143.8441 | 0.0611 | 1.0000 | |
| Stage 1 - stage 3 | 3.7268 | 3.7433 | 20.9495 | 0.9956 | 0.9961 | |
| Stage 1 - stage 4 | -31.5498 | 3.9515 | 27.6299 | -7.9844 | 0.0001 | *** |
| Stage 1 - stage 5 | 29.0889 | 4.5641 | 43.7012 | 6.3734 | 0.0001 | *** |
| Stage 1 - stage 6 | 14.4145 | 5.3614 | 72.4407 | 2.6886 | 0.2531 | |
| Stage 2 - stage 3 | -3.5126 | 4.1441 | 30.7036 | -0.8476 | 0.9992 | |
| Stage 2 - stage 4 | -31.3357 | 4.3330 | 38.5610 | -7.2319 | 0.0001 | *** |
| Stage 2 - stage 5 | -28.8748 | 4.8982 | 55.0465 | -5.8950 | 0.0001 | *** |
| Stage 2 - stage 6 | -14.2003 | 5.6485 | 82.9349 | -2.5140 | 0.3463 | |

| | | | | | | |
|---|---|---|---|---|---|---|
| Stage 3 - stage 4 | -27.8231 | 3.1299 | 154.2706 | -8.8894 | 0.0001 | *** |
| Stage 3 - stage 5 | 25.3622 | 3.9646 | 149.9901 | 6.3971 | 0.0001 | *** |
| Stage 3 - stage 6 | 10.6877 | 4.7689 | 158.9345 | 2.2411 | 0.5228 | |
| Stage 4 - stage 5 | -2.4609 | 3.9085 | 150.9554 | -0.6296 | 1.0000 | |
| Stage 4 - stage 6 | -17.1354 | 4.9327 | 154.8341 | -3.4739 | 0.0314 | * |
| Stage 5 - stage 6 | -14.6745 | 5.2982 | 148.2074 | -2.7697 | 0.2041 | |
| Stage 6 - stage 7 | -8.1667 | 5.1786 | 105.7313 | -1.5770 | 0.9691 | |
| Stage 7 - stage 8 | -3.0430 | 1.0103 | 33.1804 | -3.0120 | 0.0243 | * |
| Dopamine | estimate | SE | df | t.ratio | p.value | |
| Stage 1 - stage 2 | -0.0029 | 0.0066 | 129.5853 | -0.4429 | 1.0000 | |
| Stage 1 - stage 3 | -0.0184 | 0.0066 | 39.6540 | -2.7751 | 0.2294 | |
| Stage 1 - stage 4 | 0.0104 | 0.0073 | 52.0272 | 1.4298 | 0.9519 | |
| Stage 1 - stage 5 | 0.0370 | 0.0081 | 68.5808 | 4.5767 | 0.0012 | ** |
| Stage 1 - stage 6 | 0.0359 | 0.0101 | 98.8083 | 3.5537 | 0.0274 | * |
| Stage 2 - stage 3 | 0.0155 | 0.0070 | 47.0390 | 2.2013 | 0.5569 | |
| Stage 2 - stage 4 | 0.0075 | 0.0076 | 59.0314 | 0.9793 | 0.9976 | |
| Stage 2 - stage 5 | -0.0399 | 0.0084 | 74.5799 | -4.7376 | 0.0006 | *** |
| Stage 2 - stage 6 | -0.0388 | 0.0104 | 102.0448 | -3.7402 | 0.0152 | * |
| Stage 3 - stage 4 | -0.0080 | 0.0059 | 138.9777 | -1.3546 | 0.9699 | |
| Stage 3 - stage 5 | 0.0554 | 0.0070 | 138.9975 | 7.9619 | 0.0001 | *** |
| Stage 3 - stage 6 | 0.0543 | 0.0092 | 138.9965 | 5.9004 | 0.0001 | *** |
| Stage 4 - stage 5 | 0.0474 | 0.0074 | 132.6073 | 6.4371 | 0.0001 | *** |
| Stage 4 - stage 6 | 0.0462 | 0.0096 | 137.2635 | 4.8153 | 0.0002 | *** |
| Stage 5 - stage 6 | -0.0012 | 0.0101 | 133.7009 | -0.1168 | 1.0000 | |
| Stage 6 - stage 7 | 0.3261 | 0.0381 | 51.6950 | 8.5689 | 0.0001 | *** |
| Stage 7 - stage 8 | 0.1374 | 0.0482 | 29.4633 | 2.8533 | 0.0372 | * |
| Tyramine | estimate | SE | df | t.ratio | p.value | |
| Stage 1 - stage 2 | 0.0031 | 0.0031 | 135.1796 | 1.0269 | 0.9968 | |
| Stage 1 - stage 3 | 0.0025 | 0.0045 | 17.6703 | 0.5606 | 1.0000 | |
| Stage 1 - stage 4 | -0.0064 | 0.0046 | 19.8186 | -1.3759 | 0.9555 | |
| Stage 1 - stage 5 | 0.0129 | 0.0049 | 24.7723 | 2.6203 | 0.3205 | |

| | | | | | | |
|---|---|---|---|---|---|---|
| Stage 1 - stage 6 | 0.0070 | 0.0056 | 38.6179 | 1.2596 | 0.9795 | |
| Stage 2 - stage 3 | 0.0006 | 0.0045 | 18.2171 | 0.1342 | 1.0000 | |
| Stage 2 - stage 4 | -0.0032 | 0.0047 | 20.3970 | -0.6945 | 0.9998 | |
| Stage 2 - stage 5 | -0.0098 | 0.0050 | 25.4058 | -1.9715 | 0.7068 | |
| Stage 2 - stage 6 | -0.0039 | 0.0056 | 39.3510 | -0.6938 | 0.9999 | |
| Stage 3 - stage 4 | -0.0039 | 0.0026 | 141.6121 | -1.5086 | 0.9360 | |
| Stage 3 - stage 5 | 0.0104 | 0.0031 | 143.0217 | 3.3357 | 0.0482 | * |
| Stage 3 - stage 6 | 0.0045 | 0.0041 | 141.1092 | 1.1098 | 0.9938 | |
| Stage 4 - stage 5 | 0.0065 | 0.0031 | 136.4016 | 2.0816 | 0.6366 | |
| Stage 4 - stage 6 | 0.0006 | 0.0041 | 137.7285 | 0.1564 | 1.0000 | |
| Stage 5 - stage 6 | -0.0059 | 0.0044 | 135.9291 | -1.3477 | 0.9710 | |
| Stage 6 - stage 7 | 0.0487 | 0.0089 | 67.2207 | 5.4612 | 0.0001 | *** |
| Stage 7 - stage 8 | 0.0186 | 0.0105 | 29.9190 | 1.7722 | 0.3062 | |

| Serotonin | estimate | SE | df | t.ratio | p.value | |
|---|---|---|---|---|---|---|
| Stage 1 - stage 2 | 0.0136 | 0.0296 | 85.4927 | 0.4575 | 1.0000 | |
| Stage 1 - stage 3 | -0.0608 | 0.0477 | 29.9788 | -1.2728 | 0.9766 | |
| Stage 1 - stage 4 | 0.0420 | 0.0437 | 21.5842 | 0.9606 | 0.9972 | |
| Stage 1 - stage 5 | 0.0424 | 0.0539 | 43.4003 | 0.7865 | 0.9997 | |
| Stage 1 - stage 6 | 0.5913 | 0.0538 | 42.8065 | 10.9821 | 0.0001 | *** |
| Stage 2 - stage 3 | -0.0284 | 0.0451 | 24.1479 | -0.6307 | 0.9999 | |
| Stage 2 - stage 4 | -0.0472 | 0.0490 | 32.6983 | -0.9634 | 0.9976 | |
| Stage 2 - stage 5 | -0.0288 | 0.0550 | 45.9839 | -0.5239 | 1.0000 | |
| Stage 2 - stage 6 | -0.5777 | 0.0550 | 45.3799 | -10.5118 | 0.0001 | *** |
| Stage 3 - stage 4 | -0.0188 | 0.0310 | 90.6319 | -0.6049 | 1.0000 | |
| Stage 3 - stage 5 | 0.0004 | 0.0396 | 90.0450 | 0.0094 | 1.0000 | |
| Stage 3 - stage 6 | 0.5493 | 0.0400 | 91.9563 | 13.7432 | 0.0001 | *** |
| Stage 4 - stage 5 | -0.0184 | 0.0424 | 86.8150 | -0.4334 | 1.0000 | |
| Stage 4 - stage 6 | 0.5305 | 0.0458 | 90.9047 | 11.5921 | 0.0001 | *** |
| Stage 5 - stage 6 | 0.5489 | 0.0521 | 89.9877 | 10.5370 | 0.0001 | *** |
| Stage 6 - stage 7 | -0.2756 | 0.1189 | 90.7031 | -2.3176 | 0.6121 | |

| | | | | | | |
|---|---|---|---|---|---|---|
| Stage 7 - stage 8 | 0.3506 | 0.1697 | 23.9792 | 2.0657 | 0.1930 | |
| | | | | | | |
| Tryptophan | estimate | SE | df | t.ratio | p.value | |
| Stage 1 - stage 2 | 0.0390 | 0.3738 | 153.2773 | 0.1044 | 1.0000 | |
| Stage 1 - stage 3 | -3.2024 | 0.8449 | 15.3034 | -3.7902 | 0.0517 | * |
| Stage 1 - stage 4 | 1.3605 | 0.8287 | 14.1617 | 1.6417 | 0.8673 | |
| Stage 1 - stage 5 | 2.2181 | 0.8738 | 17.4311 | 2.5385 | 0.3782 | |
| Stage 1 - stage 6 | 2.3965 | 0.9410 | 23.1600 | 2.5467 | 0.3611 | |
| | | | | | | |
| Stage 2 - stage 3 | -1.3215 | 0.8310 | 14.3211 | -1.5903 | 0.8875 | |
| Stage 2 - stage 4 | -3.1634 | 0.8471 | 15.4687 | -3.7343 | 0.0566 | * |
| Stage 2 - stage 5 | -2.1791 | 0.8759 | 17.6061 | -2.4878 | 0.4042 | |
| Stage 2 - stage 6 | -2.3575 | 0.9430 | 23.3568 | -2.5000 | 0.3859 | |
| | | | | | | |
| Stage 3 - stage 4 | -1.8419 | 0.3311 | 155.8332 | -5.5630 | 0.0001 | *** |
| Stage 3 - stage 5 | 0.8576 | 0.4058 | 156.6401 | 2.1133 | 0.6141 | |
| Stage 3 - stage 6 | 1.0360 | 0.5350 | 155.7889 | 1.9365 | 0.7345 | |
| | | | | | | |
| Stage 4 - stage 5 | -0.9843 | 0.4194 | 153.9073 | -2.3469 | 0.4485 | |
| Stage 4 - stage 6 | -0.8059 | 0.5530 | 154.4181 | -1.4572 | 0.9495 | |
| | | | | | | |
| Stage 5 - stage 6 | 0.1784 | 0.5821 | 153.5676 | 0.3064 | 1.0000 | |
| Stage 6 - stage 7 | -1.8985 | 1.0141 | 59.7509 | -1.8721 | 0.8779 | |
| | | | | | | |
| Stage 7 - stage 8 | 1.5080 | 0.6623 | 22.0000 | 2.2769 | 0.1342 | |
| | | | | | | |
| Tryptamine | estimate | SE | df | t.ratio | p.value | |
| Stage 1 - stage 2 | 0.0109 | 0.0125 | 117.6303 | 0.8710 | 0.9993 | |
| Stage 1 - stage 3 | 0.0146 | 0.0149 | 29.3669 | 0.9817 | 0.9970 | |
| Stage 1 - stage 4 | -0.0213 | 0.0147 | 27.3582 | -1.4485 | 0.9423 | |
| Stage 1 - stage 5 | -0.0201 | 0.0163 | 39.2910 | -1.2299 | 0.9830 | |
| Stage 1 - stage 6 | -0.0159 | 0.0183 | 55.0954 | -0.8643 | 0.9992 | |
| | | | | | | |
| Stage 2 - stage 3 | 0.0322 | 0.0140 | 22.8939 | 2.3073 | 0.4987 | |
| Stage 2 - stage 4 | 0.0255 | 0.0141 | 24.7450 | 1.8049 | 0.8008 | |
| Stage 2 - stage 5 | 0.0310 | 0.0157 | 34.5698 | 1.9788 | 0.7036 | |
| Stage 2 - stage 6 | 0.0268 | 0.0177 | 50.6491 | 1.5077 | 0.9315 | |
| | | | | | | |
| Stage 3 - stage 4 | -0.0067 | 0.0092 | 121.9310 | -0.7330 | 0.9999 | |
| Stage 3 - stage 5 | 0.0012 | 0.0117 | 123.9014 | 0.1040 | 1.0000 | |
| Stage 3 - stage 6 | 0.0055 | 0.0142 | 122.1571 | 0.3841 | 1.0000 | |

| | | | | | |
|---|---|---|---|---|---|
| Stage 4 - stage 5 | -0.0055 | 0.0114 | 117.7812 | -0.4815 | 1.0000 |
| Stage 4 - stage 6 | -0.0012 | 0.0143 | 119.0582 | -0.0871 | 1.0000 |
| Stage 5 - stage 6 | 0.0042 | 0.0155 | 116.6436 | 0.2735 | 1.0000 |
| Stage 6 - stage 7 | 0.0124 | 0.0193 | 85.4670 | 0.6429 | 1.0000 |
| Stage 7 - stage 8 | 0.0061 | 0.0175 | 32.2935 | 0.3478 | 0.9853 |

**Table s6: Hawthorn race developmental stage comparisons for each neurotransmitter.**

| Histidine | estimate | SE | df | t.ratio | p.value | |
|---|---|---|---|---|---|---|
| Stage 1 - stage 2 | 9.2760 | 3.3472 | 150.2634 | 2.7713 | 0.2033 | |
| Stage 1 - stage 3 | -8.1420 | 3.6374 | 35.6718 | -2.2384 | 0.5348 | |
| Stage 1 - stage 4 | -18.5680 | 4.2215 | 57.4697 | -4.3984 | 0.0026 | ** |
| Stage 1 - stage 5 | 1.4017 | 4.9156 | 79.2112 | 0.2852 | 1.0000 | |
| Stage 1 - stage 6 | -4.5546 | 4.6661 | 62.2434 | -0.9761 | 0.9977 | |
| Stage 2 - stage 3 | 17.4180 | 3.4578 | 29.7771 | 5.0373 | 0.0011 | ** |
| Stage 2 - stage 4 | -9.2919 | 4.0678 | 51.2306 | -2.2843 | 0.5002 | |
| Stage 2 - stage 5 | 7.8743 | 4.7842 | 73.8749 | 1.6459 | 0.8858 | |
| Stage 2 - stage 6 | 13.8306 | 4.5274 | 56.8971 | 3.0548 | 0.1196 | |
| Stage 3 - stage 4 | -26.7099 | 3.2232 | 160.6241 | -8.2867 | 0.0001 | *** |
| Stage 3 - stage 5 | 9.5437 | 4.1790 | 155.8926 | 2.2837 | 0.4926 | |
| Stage 3 - stage 6 | 3.5874 | 3.9071 | 134.5693 | 0.9182 | 0.9988 | |
| Stage 4 - stage 5 | -17.1662 | 4.4386 | 154.5738 | -3.8675 | 0.0086 | ** |
| Stage 4 - stage 6 | -23.1225 | 4.1837 | 160.9020 | -5.5268 | 0.0001 | *** |
| Stage 5 - stage 6 | -5.9563 | 4.7049 | 152.9116 | -1.2660 | 0.9821 | |
| Stage 6 - stage 7 | 7.4986 | 4.6068 | 75.3566 | 1.6277 | 0.9579 | |
| Stage 7 - stage 8 | -9.4681 | 1.9680 | 35.4192 | -4.8110 | 0.0002 | *** |
| **Serine** | **estimate** | **SE** | **df** | **t.ratio** | **p.value** | |
| Stage 1 - stage 2 | -1.4183 | 1.8679 | 141.2302 | -0.7593 | 0.9998 | |
| Stage 1 - stage 3 | -5.3565 | 2.1221 | 23.4911 | -2.5242 | 0.3726 | |
| Stage 1 - stage 4 | -10.0340 | 2.4487 | 39.8930 | -4.0976 | 0.0094 | ** |
| Stage 1 - stage 5 | 13.0445 | 2.7836 | 53.8388 | 4.6863 | 0.0011 | ** |
| Stage 1 - stage 6 | 0.2133 | 2.7264 | 46.6659 | 0.0782 | 1.0000 | |
| Stage 2 - stage 3 | 3.9382 | 2.0498 | 20.8278 | 1.9212 | 0.7351 | |
| Stage 2 - stage 4 | -11.4523 | 2.3864 | 36.8641 | -4.7990 | 0.0014 | ** |
| Stage 2 - stage 5 | -14.4628 | 2.7289 | 51.0300 | -5.2999 | 0.0001 | *** |
| Stage 2 - stage 6 | -1.6316 | 2.6706 | 43.9618 | -0.6110 | 1.0000 | |
| Stage 3 - stage 4 | -15.3905 | 1.8540 | 152.4109 | -8.3011 | 0.0001 | *** |
| Stage 3 - stage 5 | 18.4009 | 2.3417 | 144.0862 | 7.8580 | 0.0001 | *** |
| Stage 3 - stage 6 | 5.5698 | 2.2836 | 129.6390 | 2.4391 | 0.3880 | |

| | | | | | | |
|---|---|---|---|---|---|---|
| Stage 4 - stage 5 | 3.0104 | 2.4795 | 144.9630 | 1.2141 | 0.9871 | |
| Stage 4 - stage 6 | -9.8207 | 2.4253 | 150.4528 | -4.0493 | 0.0045 | ** |
| Stage 5 - stage 6 | -12.8311 | 2.6298 | 139.1037 | -4.8790 | 0.0002 | *** |
| Stage 6 - stage 7 | 9.5102 | 3.0334 | 54.4169 | 3.1352 | 0.1522 | |
| Stage 7 - stage 8 | -5.6389 | 2.1560 | 34.5884 | -2.6154 | 0.0603 | |
| Histamine | estimate | SE | df | t.ratio | p.value | |
| Stage 1 - stage 2 | 0.0030 | 0.0045 | 121.9995 | 0.6518 | 1.0000 | |
| Stage 1 - stage 3 | 0.0059 | 0.0068 | 16.9106 | 0.8617 | 0.9987 | |
| Stage 1 - stage 4 | -0.0244 | 0.0071 | 20.3297 | -3.4219 | 0.0818 | |
| Stage 1 - stage 5 | 0.0230 | 0.0077 | 26.6557 | 2.9844 | 0.1684 | |
| Stage 1 - stage 6 | 0.0986 | 0.0077 | 25.6194 | 12.8708 | 0.0001 | *** |
| Stage 2 - stage 3 | -0.0029 | 0.0068 | 16.9128 | -0.4270 | 1.0000 | |
| Stage 2 - stage 4 | -0.0215 | 0.0071 | 20.3318 | -3.0085 | 0.1759 | |
| Stage 2 - stage 5 | -0.0201 | 0.0077 | 26.6575 | -2.6018 | 0.3267 | |
| Stage 2 - stage 6 | -0.0956 | 0.0077 | 25.6213 | -12.4848 | 0.0001 | *** |
| Stage 3 - stage 4 | -0.0186 | 0.0041 | 128.2996 | -4.5462 | 0.0007 | *** |
| Stage 3 - stage 5 | 0.0172 | 0.0052 | 130.5207 | 3.3064 | 0.0532 | |
| Stage 3 - stage 6 | 0.0927 | 0.0051 | 131.9887 | 18.0226 | 0.0001 | *** |
| Stage 4 - stage 5 | -0.0014 | 0.0051 | 123.1360 | -0.2740 | 1.0000 | |
| Stage 4 - stage 6 | 0.0741 | 0.0051 | 125.2129 | 14.5311 | 0.0001 | *** |
| Stage 5 - stage 6 | 0.0755 | 0.0056 | 122.7340 | 13.5327 | 0.0001 | *** |
| Stage 6 - stage 7 | 0.4911 | 0.0325 | 42.5499 | 15.1207 | 0.0001 | *** |
| Stage 7 - stage 8 | 0.2389 | 0.0455 | 34.8184 | 5.2453 | 0.0001 | *** |
| Aspartate | estimate | SE | df | t.ratio | p.value | |
| Stage 1 - stage 2 | -1.4488 | 3.7653 | 148.7253 | -0.3848 | 1.0000 | |
| Stage 1 - stage 3 | 2.6594 | 7.0984 | 13.3624 | 0.3747 | 1.0000 | |
| Stage 1 - stage 4 | -51.1903 | 7.4971 | 16.5096 | -6.8280 | 0.0002 | *** |
| Stage 1 - stage 5 | 30.0268 | 7.9971 | 20.9307 | 3.7547 | 0.0409 | * |
| Stage 1 - stage 6 | 25.4946 | 7.9695 | 20.4891 | 3.1990 | 0.1244 | |
| Stage 2 - stage 3 | -4.1083 | 7.0651 | 13.1084 | -0.5815 | 1.0000 | |
| Stage 2 - stage 4 | -52.6391 | 7.4656 | 16.2307 | -7.0509 | 0.0001 | *** |

| | | | | | | |
|---|---|---|---|---|---|---|
| Stage 2 - stage 5 | -31.4756 | 7.9675 | 20.6243 | -3.9505 | 0.0275 | * |
| Stage 2 - stage 6 | -26.9434 | 7.9398 | 20.1867 | -3.3934 | 0.0868 | |
| | | | | | | |
| Stage 3 - stage 4 | -48.5308 | 3.7572 | 158.0238 | -12.9167 | 0.0001 | *** |
| Stage 3 - stage 5 | 27.3673 | 4.8379 | 159.3046 | 5.6568 | 0.0001 | *** |
| Stage 3 - stage 6 | 22.8352 | 4.8331 | 160.4201 | 4.7247 | 0.0003 | *** |
| | | | | | | |
| Stage 4 - stage 5 | -21.1635 | 4.7285 | 149.9532 | -4.4757 | 0.0009 | *** |
| Stage 4 - stage 6 | -25.6957 | 4.7262 | 152.1083 | -5.4369 | 0.0001 | *** |
| | | | | | | |
| Stage 5 - stage 6 | -4.5322 | 5.1331 | 149.2986 | -0.8829 | 0.9992 | |
| Stage 6 - stage 7 | 138.0729 | 14.7200 | 23.1542 | 9.3800 | 0.0001 | *** |
| | | | | | | |
| Stage 7 - stage 8 | 1.9889 | 11.9196 | 34.6003 | 0.1669 | 0.9983 | |
| | | | | | | |
| Glutamate | estimate | SE | df | t.ratio | p.value | |
| Stage 1 - stage 2 | 10.2681 | 7.1871 | 150.0784 | 1.4287 | 0.9560 | |
| Stage 1 - stage 3 | -9.7917 | 9.4441 | 20.5562 | -1.0368 | 0.9945 | |
| Stage 1 - stage 4 | -54.0539 | 10.4326 | 29.4495 | -5.1813 | 0.0008 | *** |
| Stage 1 - stage 5 | -4.7870 | 11.5635 | 40.6643 | -0.4140 | 1.0000 | |
| Stage 1 - stage 6 | -5.9932 | 11.4700 | 38.2213 | -0.5225 | 1.0000 | |
| | | | | | | |
| Stage 2 - stage 3 | 20.0598 | 9.4475 | 20.5660 | 2.1233 | 0.6138 | |
| Stage 2 - stage 4 | -43.7859 | 10.4356 | 29.4562 | -4.1958 | 0.0102 | * |
| Stage 2 - stage 5 | 15.0551 | 11.5662 | 40.6661 | 1.3016 | 0.9742 | |
| Stage 2 - stage 6 | 16.2612 | 11.4728 | 38.2242 | 1.4174 | 0.9527 | |
| | | | | | | |
| Stage 3 - stage 4 | -63.8456 | 6.7773 | 160.6880 | -9.4205 | 0.0001 | *** |
| Stage 3 - stage 5 | 5.0047 | 8.6549 | 160.9555 | 0.5782 | 1.0000 | |
| Stage 3 - stage 6 | 3.7985 | 8.5995 | 158.9816 | 0.4417 | 1.0000 | |
| | | | | | | |
| Stage 4 - stage 5 | -58.8410 | 8.8143 | 151.6863 | -6.6757 | 0.0001 | *** |
| Stage 4 - stage 6 | -60.0471 | 8.7713 | 155.7959 | -6.8459 | 0.0001 | *** |
| | | | | | | |
| Stage 5 - stage 6 | -1.2061 | 9.5939 | 151.2755 | -0.1257 | 1.0000 | |
| Stage 6 - stage 7 | 130.7025 | 13.1765 | 45.8484 | 9.9194 | 0.0001 | *** |
| | | | | | | |
| Stage 7 - stage 8 | 0.5994 | 10.5082 | 32.8034 | 0.0570 | 0.9999 | |
| | | | | | | |
| GABA | estimate | SE | df | t.ratio | p.value | |
| Stage 1 - stage 2 | 0.4085 | 0.6457 | 146.4839 | 0.6327 | 1.0000 | |

| | | | | | | |
|---|---|---|---|---|---|---|
| Stage 1 - stage 3 | -0.1254 | 0.6311 | 42.5976 | -0.1987 | 1.0000 | |
| Stage 1 - stage 4 | 2.1235 | 0.7393 | 66.2943 | 2.8722 | 0.1752 | |
| Stage 1 - stage 5 | 4.5432 | 0.8546 | 82.2229 | 5.3160 | 0.0001 | *** |
| Stage 1 - stage 6 | 14.0671 | 0.8636 | 75.1907 | 16.2892 | 0.0001 | *** |
| | | | | | | |
| Stage 2 - stage 3 | 0.5339 | 0.6300 | 42.8572 | 0.8475 | 0.9993 | |
| Stage 2 - stage 4 | 2.5320 | 0.7384 | 66.7022 | 3.4292 | 0.0446 | * |
| Stage 2 - stage 5 | -4.1347 | 0.8538 | 82.6405 | -4.8428 | 0.0004 | *** |
| Stage 2 - stage 6 | -13.6586 | 0.8628 | 75.5517 | -15.8314 | 0.0001 | *** |
| | | | | | | |
| Stage 3 - stage 4 | 1.9981 | 0.5893 | 152.0020 | 3.3909 | 0.0405 | * |
| Stage 3 - stage 5 | 4.6685 | 0.7418 | 138.7799 | 6.2935 | 0.0001 | *** |
| Stage 3 - stage 6 | 14.1924 | 0.7563 | 120.7116 | 18.7656 | 0.0001 | *** |
| | | | | | | |
| Stage 4 - stage 5 | 6.6667 | 0.7915 | 152.1419 | 8.4226 | 0.0001 | *** |
| Stage 4 - stage 6 | 16.1906 | 0.8059 | 157.6172 | 20.0903 | 0.0001 | *** |
| | | | | | | |
| Stage 5 - stage 6 | 9.5239 | 0.8866 | 150.5125 | 10.7419 | 0.0001 | *** |
| Stage 6 - stage 7 | 44.5772 | 2.8134 | 31.8033 | 15.8444 | 0.0001 | *** |
| | | | | | | |
| Stage 7 - stage 8 | 0.1193 | 3.1720 | 32.6319 | 0.0376 | 1.0000 | |
| | | | | | | |
| Dopa | estimate | SE | df | t.ratio | p.value | |
| Stage 1 - stage 2 | -0.0152 | 0.0260 | 149.2417 | -0.5864 | 1.0000 | |
| Stage 1 - stage 3 | -0.0574 | 0.0259 | 50.5113 | -2.2141 | 0.5476 | |
| Stage 1 - stage 4 | -0.1969 | 0.0317 | 84.6373 | -6.2059 | 0.0001 | *** |
| Stage 1 - stage 5 | -0.0065 | 0.0353 | 90.9806 | -0.1840 | 1.0000 | |
| Stage 1 - stage 6 | -0.0008 | 0.0356 | 85.1021 | -0.0212 | 1.0000 | |
| | | | | | | |
| Stage 2 - stage 3 | 0.0421 | 0.0249 | 45.0092 | 1.6911 | 0.8628 | |
| Stage 2 - stage 4 | -0.2121 | 0.0309 | 80.2057 | -6.8597 | 0.0001 | *** |
| Stage 2 - stage 5 | -0.0087 | 0.0346 | 87.4519 | -0.2521 | 1.0000 | |
| Stage 2 - stage 6 | -0.0145 | 0.0349 | 81.6114 | -0.4143 | 1.0000 | |
| | | | | | | |
| Stage 3 - stage 4 | -0.2542 | 0.0258 | 154.8789 | -9.8450 | 0.0001 | *** |
| Stage 3 - stage 5 | 0.0509 | 0.0307 | 137.2628 | 1.6545 | 0.8851 | |
| Stage 3 - stage 6 | 0.0566 | 0.0313 | 123.8106 | 1.8118 | 0.8085 | |
| | | | | | | |
| Stage 4 - stage 5 | -0.2034 | 0.0341 | 153.5463 | -5.9646 | 0.0001 | *** |
| Stage 4 - stage 6 | -0.1976 | 0.0345 | 157.0370 | -5.7267 | 0.0001 | *** |

| | | | | | | |
|---|---|---|---|---|---|---|
| Stage 5 - stage 6 | 0.0057 | 0.0368 | 151.8266 | 0.1559 | 1.0000 | |
| Stage 6 - stage 7 | -0.0487 | 0.0338 | 105.8274 | -1.4395 | 0.9865 | |
| Stage 7 - stage 8 | -0.0060 | 0.0017 | 33.1420 | -3.5825 | 0.0057 | ** |
| Octopamine | estimate | SE | df | t.ratio | p.value | |
| Stage 1 - stage 2 | 0.0006 | 0.0447 | 78.2365 | 0.0131 | 1.0000 | |
| Stage 1 - stage 3 | 0.0139 | 0.0604 | 16.4742 | 0.2296 | 1.0000 | |
| Stage 1 - stage 4 | -0.0067 | 0.0600 | 16.1625 | -0.1125 | 1.0000 | |
| Stage 1 - stage 5 | 0.0112 | 0.0653 | 21.5997 | 0.1722 | 1.0000 | |
| Stage 1 - stage 6 | 0.0916 | 0.0636 | 19.3057 | 1.4405 | 0.9400 | |
| Stage 2 - stage 3 | -0.0133 | 0.0568 | 13.2860 | -0.2337 | 1.0000 | |
| Stage 2 - stage 4 | -0.0107 | 0.0620 | 18.1907 | -0.1718 | 1.0000 | |
| Stage 2 - stage 5 | -0.0910 | 0.0602 | 16.0158 | -1.5113 | 0.9168 | |
| Stage 2 - stage 6 | -0.0062 | 0.0564 | 12.9841 | -0.1092 | 1.0000 | |
| Stage 3 - stage 4 | 0.0071 | 0.0370 | 81.0528 | 0.1925 | 1.0000 | |
| Stage 3 - stage 5 | -0.0026 | 0.0463 | 84.3525 | -0.0565 | 1.0000 | |
| Stage 3 - stage 6 | 0.0777 | 0.0446 | 86.9104 | 1.7414 | 0.8437 | |
| Stage 4 - stage 5 | 0.0045 | 0.0434 | 79.3577 | 0.1037 | 1.0000 | |
| Stage 4 - stage 6 | 0.0848 | 0.0420 | 82.5509 | 2.0209 | 0.6781 | |
| Stage 5 - stage 6 | 0.0803 | 0.0478 | 81.8784 | 1.6801 | 0.8719 | |
| Stage 6 - stage 7 | 0.2462 | 0.0482 | 37.7643 | 5.1098 | 0.0009 | *** |
| Stage 7 - stage 8 | 0.1023 | 0.0294 | 34.0988 | 3.4782 | 0.0073 | ** |
| Tyrosine | estimate | SE | df | t.ratio | p.value | |
| Stage 1 - stage 2 | -0.1615 | 3.2025 | 143.9551 | -0.0504 | 1.0000 | |
| Stage 1 - stage 3 | 0.8927 | 3.6657 | 20.3734 | 0.2435 | 1.0000 | |
| Stage 1 - stage 4 | -59.8429 | 4.2644 | 33.6500 | -14.0332 | 0.0001 | *** |
| Stage 1 - stage 5 | 41.4830 | 4.7469 | 44.5304 | 8.7390 | 0.0001 | *** |
| Stage 1 - stage 6 | 32.0165 | 4.6897 | 39.2125 | 6.8270 | 0.0001 | *** |
| Stage 2 - stage 3 | -1.0542 | 3.6144 | 19.2515 | -0.2917 | 1.0000 | |
| Stage 2 - stage 4 | -60.0044 | 4.2204 | 32.3473 | -14.2178 | 0.0001 | *** |
| Stage 2 - stage 5 | -41.6445 | 4.7074 | 43.2119 | -8.8467 | 0.0001 | *** |
| Stage 2 - stage 6 | -32.1780 | 4.6497 | 37.9893 | -6.9204 | 0.0001 | *** |

| | | | | | | |
|---|---|---|---|---|---|---|
| Stage 3 - stage 4 | -58.9502 | 3.2072 | 149.0144 | -18.3804 | 0.0001 | *** |
| Stage 3 - stage 5 | 40.5903 | 3.9062 | 143.0938 | 10.3914 | 0.0001 | *** |
| Stage 3 - stage 6 | 31.1238 | 3.8641 | 123.5099 | 8.0546 | 0.0001 | *** |
| Stage 4 - stage 5 | -18.3599 | 4.0528 | 146.7099 | -4.5302 | 0.0007 | *** |
| Stage 4 - stage 6 | -27.8264 | 4.0274 | 155.4292 | -6.9092 | 0.0001 | *** |
| Stage 5 - stage 6 | -9.4665 | 4.3606 | 147.8187 | -2.1709 | 0.5731 | |
| Stage 6 - stage 7 | -25.1479 | 4.4380 | 57.7244 | -5.6665 | 0.0001 | *** |
| Stage 7 - stage 8 | -5.0456 | 1.0150 | 34.7425 | -4.9710 | 0.0001 | *** |
| Dopamine | estimate | SE | df | t.ratio | p.value | |
| Stage 1 - stage 2 | 0.0179 | 0.0064 | 130.3156 | 2.7912 | 0.1962 | |
| Stage 1 - stage 3 | -0.0119 | 0.0070 | 43.8055 | -1.6919 | 0.8622 | |
| Stage 1 - stage 4 | 0.0030 | 0.0079 | 59.5628 | 0.3819 | 1.0000 | |
| Stage 1 - stage 5 | 0.0026 | 0.0091 | 76.0666 | 0.2900 | 1.0000 | |
| Stage 1 - stage 6 | 0.0243 | 0.0091 | 73.0937 | 2.6703 | 0.2620 | |
| Stage 2 - stage 3 | 0.0297 | 0.0070 | 43.9311 | 4.2419 | 0.0056 | ** |
| Stage 2 - stage 4 | 0.0209 | 0.0078 | 59.7562 | 2.6598 | 0.2716 | |
| Stage 2 - stage 5 | 0.0152 | 0.0091 | 76.2884 | 1.6825 | 0.8705 | |
| Stage 2 - stage 6 | -0.0064 | 0.0091 | 73.3004 | -0.7074 | 0.9999 | |
| Stage 3 - stage 4 | -0.0089 | 0.0066 | 138.7670 | -1.3402 | 0.9722 | |
| Stage 3 - stage 5 | 0.0145 | 0.0082 | 136.8448 | 1.7644 | 0.8341 | |
| Stage 3 - stage 6 | 0.0362 | 0.0083 | 133.0699 | 4.3522 | 0.0015 | ** |
| Stage 4 - stage 5 | 0.0056 | 0.0085 | 132.1090 | 0.6639 | 0.9999 | |
| Stage 4 - stage 6 | 0.0273 | 0.0086 | 136.8735 | 3.1653 | 0.0782 | |
| Stage 5 - stage 6 | 0.0217 | 0.0095 | 132.5954 | 2.2841 | 0.4930 | |
| Stage 6 - stage 7 | 0.4249 | 0.0366 | 43.3556 | 11.6190 | 0.0001 | *** |
| Stage 7 - stage 8 | -0.0389 | 0.0497 | 29.5559 | -0.7835 | 0.8613 | |
| Tyramine | estimate | SE | df | t.ratio | p.value | |
| Stage 1 - stage 2 | 0.0004 | 0.0030 | 135.9730 | 0.1404 | 1.0000 | |
| Stage 1 - stage 3 | 0.0034 | 0.0045 | 17.5711 | 0.7463 | 0.9997 | |
| Stage 1 - stage 4 | -0.0255 | 0.0048 | 22.7596 | -5.2865 | 0.0011 | ** |
| Stage 1 - stage 5 | 0.0215 | 0.0052 | 28.9461 | 4.1524 | 0.0116 | * |

| | | | | | | |
|---|---|---|---|---|---|---|
| Stage 1 - stage 6 | 0.0242 | 0.0053 | 30.0147 | 4.5980 | 0.0035 | ** |
| Stage 2 - stage 3 | -0.0029 | 0.0044 | 15.5702 | -0.6743 | 0.9999 | |
| Stage 2 - stage 4 | -0.0251 | 0.0047 | 20.5446 | -5.3425 | 0.0013 | ** |
| Stage 2 - stage 5 | -0.0211 | 0.0051 | 26.5769 | -4.1680 | 0.0124 | * |
| Stage 2 - stage 6 | -0.0238 | 0.0051 | 27.6413 | -4.6222 | 0.0038 | ** |
| Stage 3 - stage 4 | -0.0221 | 0.0027 | 143.6473 | -8.1276 | 0.0001 | *** |
| Stage 3 - stage 5 | 0.0182 | 0.0034 | 146.1464 | 5.2853 | 0.0001 | *** |
| Stage 3 - stage 6 | 0.0208 | 0.0036 | 146.8018 | 5.8266 | 0.0001 | *** |
| Stage 4 - stage 5 | -0.0040 | 0.0035 | 137.0306 | -1.1506 | 0.9916 | |
| Stage 4 - stage 6 | -0.0013 | 0.0036 | 139.6144 | -0.3674 | 1.0000 | |
| Stage 5 - stage 6 | 0.0027 | 0.0038 | 136.6062 | 0.6924 | 0.9999 | |
| Stage 6 - stage 7 | 0.0214 | 0.0080 | 46.5211 | 2.6795 | 0.3735 | |
| Stage 7 - stage 8 | -0.0045 | 0.0104 | 31.4902 | -0.4360 | 0.9718 | |

| Serotonin | estimate | SE | df | t.ratio | p.value | |
|---|---|---|---|---|---|---|
| Stage 1 - stage 2 | 0.0060 | 0.0273 | 89.8211 | 0.2209 | 1.0000 | |
| Stage 1 - stage 3 | 0.0562 | 0.0426 | 20.5010 | 1.3176 | 0.9669 | |
| Stage 1 - stage 4 | -0.0151 | 0.0496 | 31.2669 | -0.3041 | 1.0000 | |
| Stage 1 - stage 5 | -0.0227 | 0.0508 | 36.3307 | -0.4476 | 1.0000 | |
| Stage 1 - stage 6 | 0.1396 | 0.0503 | 31.7630 | 2.7759 | 0.2376 | |
| Stage 2 - stage 3 | -0.0502 | 0.0435 | 21.1722 | -1.1519 | 0.9876 | |
| Stage 2 - stage 4 | -0.0091 | 0.0504 | 31.6932 | -0.1799 | 1.0000 | |
| Stage 2 - stage 5 | 0.0288 | 0.0516 | 36.6551 | 0.5577 | 1.0000 | |
| Stage 2 - stage 6 | -0.1335 | 0.0510 | 32.1716 | -2.6162 | 0.3121 | |
| Stage 3 - stage 4 | 0.0411 | 0.0383 | 92.3420 | 1.0737 | 0.9951 | |
| Stage 3 - stage 5 | -0.0789 | 0.0405 | 94.9835 | -1.9471 | 0.7267 | |
| Stage 3 - stage 6 | 0.0834 | 0.0404 | 91.1039 | 2.0635 | 0.6493 | |
| Stage 4 - stage 5 | -0.0378 | 0.0419 | 89.7910 | -0.9034 | 0.9989 | |
| Stage 4 - stage 6 | 0.1245 | 0.0465 | 93.0608 | 2.6755 | 0.2551 | |
| Stage 5 - stage 6 | 0.1623 | 0.0449 | 94.8561 | 3.6159 | 0.0230 | * |
| Stage 6 - stage 7 | 0.2767 | 0.1106 | 64.5195 | 2.5028 | 0.4824 | |

| | | | | | | |
|---|---|---|---|---|---|---|
| Stage 7 - stage 8 | -0.0828 | 0.1657 | 24.8909 | -0.4998 | 0.9584 | |
| | | | | | | |
| Tryptophan | estimate | SE | df | t.ratio | p.value | |
| Stage 1 - stage 2 | -0.0291 | 0.3596 | 153.2724 | -0.0808 | 1.0000 | |
| Stage 1 - stage 3 | 0.7692 | 0.8231 | 13.7971 | 0.9345 | 0.9971 | |
| Stage 1 - stage 4 | -4.6197 | 0.8570 | 16.1579 | -5.3908 | 0.0024 | ** |
| Stage 1 - stage 5 | 1.9547 | 0.8934 | 18.9130 | 2.1878 | 0.5752 | |
| Stage 1 - stage 6 | 1.1053 | 0.9029 | 19.6164 | 1.2242 | 0.9800 | |
| | | | | | | |
| Stage 2 - stage 3 | -0.7982 | 0.8205 | 13.6208 | -0.9729 | 0.9959 | |
| Stage 2 - stage 4 | -4.6487 | 0.8544 | 15.9689 | -5.4408 | 0.0022 | ** |
| Stage 2 - stage 5 | -1.9837 | 0.8910 | 18.7114 | -2.2264 | 0.5520 | |
| Stage 2 - stage 6 | -1.1344 | 0.9005 | 19.4122 | -1.2597 | 0.9753 | |
| | | | | | | |
| Stage 3 - stage 4 | -3.8505 | 0.3574 | 157.1249 | -10.7744 | 0.0001 | *** |
| Stage 3 - stage 5 | 1.1855 | 0.4544 | 158.8217 | 2.6090 | 0.2837 | |
| Stage 3 - stage 6 | 0.3362 | 0.4766 | 159.9671 | 0.7053 | 0.9999 | |
| | | | | | | |
| Stage 4 - stage 5 | -2.6650 | 0.4616 | 154.0365 | -5.7735 | 0.0001 | *** |
| Stage 4 - stage 6 | -3.5143 | 0.4825 | 155.3909 | -7.2829 | 0.0001 | *** |
| | | | | | | |
| Stage 5 - stage 6 | -0.8493 | 0.5133 | 153.9906 | -1.6546 | 0.8854 | |
| Stage 6 - stage 7 | 0.2382 | 0.8662 | 33.6721 | 0.2749 | 1.0000 | |
| | | | | | | |
| Stage 7 - stage 8 | -1.0817 | 0.6399 | 22.0000 | -1.6905 | 0.3522 | |
| | | | | | | |
| Tryptamine | estimate | SE | df | t.ratio | p.value | |
| Stage 1 - stage 2 | -0.0121 | 0.0119 | 115.5362 | -1.0167 | 0.9970 | |
| Stage 1 - stage 3 | -0.0372 | 0.0140 | 23.2468 | -2.6650 | 0.3017 | |
| Stage 1 - stage 4 | 0.0256 | 0.0148 | 28.2896 | 1.7315 | 0.8391 | |
| Stage 1 - stage 5 | -0.0396 | 0.0166 | 39.9137 | -2.3893 | 0.4358 | |
| Stage 1 - stage 6 | -0.0389 | 0.0160 | 34.4596 | -2.4229 | 0.4184 | |
| | | | | | | |
| Stage 2 - stage 3 | 0.0251 | 0.0137 | 21.5369 | 1.8361 | 0.7826 | |
| Stage 2 - stage 4 | 0.0135 | 0.0145 | 26.4791 | 0.9301 | 0.9980 | |
| Stage 2 - stage 5 | 0.0275 | 0.0163 | 38.0354 | 1.6838 | 0.8644 | |
| Stage 2 - stage 6 | 0.0268 | 0.0158 | 32.6616 | 1.6954 | 0.8576 | |
| | | | | | | |
| Stage 3 - stage 4 | -0.0116 | 0.0094 | 123.7577 | -1.2323 | 0.9853 | |
| Stage 3 - stage 5 | -0.0024 | 0.0124 | 123.9744 | -0.1916 | 1.0000 | |
| Stage 3 - stage 6 | -0.0017 | 0.0118 | 120.8547 | -0.1407 | 1.0000 | |

| | | | | | | |
|---|---|---|---|---|---|---|
| Stage 4 - stage 5 | -0.0140 | 0.0123 | 116.6200 | -1.1395 | 0.9921 | |
| Stage 4 - stage 6 | -0.0133 | 0.0117 | 120.6759 | -1.1403 | 0.9921 | |
| Stage 5 - stage 6 | 0.0007 | 0.0132 | 115.8549 | 0.0537 | 1.0000 | |
| Stage 6 - stage 7 | 0.0738 | 0.0166 | 52.3531 | 4.4421 | 0.0042 | ** |
| Stage 7 - stage 8 | -0.0740 | 0.0167 | 34.5264 | -4.4230 | 0.0005 | *** |

**Table s7: Two-way ANOVA result table showing host race comparisons at stage 4 for selected neurotransmitters.**

| Tyrosine | df | F | P | error |
|---|---|---|---|---|
| Hostrace | 1 | 13.231 | 0.005 | |
| Days | 12 | 4.936 | 0.008 | |
| Hostrace*Days | 12 | 10.978 | 0.0001 | 10 |
| | | | | |
| Tyramine | df | F | P | error |
| Hostrace | 1 | 9.705 | 0.011 | |
| Days | 12 | 3.153 | 0.039 | |
| Hostrace*Days | 12 | 4.957 | 0.008 | 10 |
| | | | | |
| DOPA | df | F | P | error |
| Hostrace | 1 | 0.649 | 0.447 | |
| Days | 12 | 0.639 | 0.764 | |
| Hostrace*Days | 12 | 0.828 | 0.632 | 7 |
| | | | | |
| Aspartate | df | F | P | error |
| Hostrace | 1 | 2.082 | 0.177 | |
| Days | 12 | 4.118 | 0.013 | |
| Hostrace*Days | 12 | 10.129 | 0.0001 | 11 |
| | | | | |
| Glutamate | df | F | P | error |
| Hostrace | 1 | 2.476 | 0.144 | |
| Days | 12 | 2.918 | 0.043 | |
| Hostrace*Days | 12 | 6.479 | 0.002 | 11 |
| | | | | |
| Tryptophan | df | F | P | error |
| Hostrace | 1 | 1.517 | 0.246 | |
| Days | 12 | 2.285 | 0.1 | |
| Hostrace*Days | 12 | 4.359 | 0.013 | 10 |

**Table s8: Two-way ANOVA result table showing host race comparisons at stage 4 for selected neurotransmitters after removing early time point till day20.**

| Tyrosine | df | F | P | error |
|---|---|---|---|---|
| Hostrace | 1 | 67.113 | 0.0001 | |
| Days | 8 | 5.786 | 0.008 | |
| Hostrace*Days | 8 | 38.149 | 0.0001 | 9 |
| | | | | |
| Tyramine | df | F | P | error |
| Hostrace | 1 | 9.261 | 0.012 | |
| Days | 8 | 2.052 | 0.142 | |
| Hostrace*Days | 7 | 4.463 | 0.017 | 10 |
| | | | | |
| DOPA | df | F | P | error |
| Hostrace | 1 | 0.4 | 0.545 | |
| Days | 8 | 0.198 | 0.983 | |
| Hostrace*Days | 7 | 0.846 | 0.581 | 8 |
| | | | | |
| Aspartate | df | F | P | error |
| Hostrace | 1 | 5.087 | 0.048 | |
| Days | 8 | 0.863 | 0.574 | |
| Hostrace*Days | 8 | 14.25 | 0.0001 | 10 |
| | | | | |
| Glutamate | df | F | P | error |
| Hostrace | 1 | 26.94 | 0.001 | |
| Days | 8 | 4.382 | 0.02 | |
| Hostrace*Days | 8 | 25.367 | 0.0001 | 9 |
| | | | | |
| Tryptophan | df | F | P | error |
| Hostrace | 1 | 1.272 | 0.286 | |
| Days | 8 | 1.716 | 0.209 | |
| Hostrace*Days | 7 | 7.193 | 0.003 | 10 |

**Table s9: Instrument method details for liquid chromatography (LC).**

| | |
|---|---|
| **Instrument** | Agilent 1290 Infinity UHPLC |
| **Column** | Waters Acquity HSS C18, 1.8u, 100mmx2.1mm |
| **Mobile Phase A** | 10mM Ammonium Acetate in Water (0.1%FA) |
| **Mobile Phase B** | Acetonitrile (0.1%FA) |
| **Flow Rate** | 0.2ml/min |
| **Column Oven** | 40°C |
| **Auto-sampler Temp.** | 10°C |
| **Injection Volume** | 10ul |
| **Run Time** | 35mins |
| **Gradient** | 0-3mins:2%B,<br>3-20mins:2-20%B,<br>20-25mins:35%B,<br>25.1mins:80%B,<br>27mins:80%B,<br>27.1-35:2%B |

**Table s10: Instrument method details for mass spctrometry (MS).**

| | |
|---|---|
| **Instrument** | Thermo Fisher- TSQ Vantage |
| **Spray Voltage (+ve)** | 3700V |
| **Vaporizer temp** | 80°C |
| **Sheath gas flow rate** | 20Arb |
| **Aux gas flow rate** | 10Arb |
| **Peak width settings** | 0.70FWHM |
| **Scan time** | 0.05s |
| **Injector settings** | 0-3mins:waste, 3-28.5mins:load, 28.5-35mins:waste |

**Table s11**: Internal standard molecular weights (MW) and expected retention times (RT) for the 14 neurotransmitters and their internal standards used for quantification in mass spectrometry.

| compound | MW | RT |
|---|---|---|
| Histidine-D3 | 329 | 7.53 |
| Histidine | 326.11 | 7.56 |
| serine-D3 | 279.2 | 8.53 |
| Serine | 276.1 | 8.59 |
| Histamine-D4 | 286.37 | 9.33 |
| Histamine | 282.37 | 9.36 |
| Aspartate-D3 | 307.1 | 9.62 |
| Aspartate | 304.1 | 9.68 |
| Glutamate-D5 | 323.26 | 10.48 |
| Glutamate | 318.1 | 10.54 |
| GABA-D6 | 280.32 | 12.96 |
| GABA | 274.28 | 13.06 |
| Dopa-D3 | 371.37 | 15.29 |
| Dopa | 368.35 | 15.33 |
| Octopamine-D3 | 327.17 | 15.44 |
| Octopamine | 324.17 | 15.5 |
| Tyrosine-D7 | 359.4 | 17.25 |
| Tyrosine | 352.2 | 17.35 |
| Dopamine-D4 | 328.37 | 17.85 |
| Dopamine | 324.3 | 17.94 |
| Tyramine-D4 | 312.36 | 20.18 |
| Tyramine | 308.36 | 20.37 |
| Serotonin-D4 | 351.4 | 22.03 |
| Serotonin | 347.38 | 22.11 |
| Tryptophan-D5 | 380.43 | 23.83 |
| Tryptophan | 375.19 | 23.91 |
| Tryptamine-D4 | 335.4 | 26.19 |
| Tryptamine | 331.38 | 26.25 |

**Table s12: Standard solution preparation concentrations for all neurotransmitters. ***

| | Standards | | | Internal standards |
|---|---|---|---|---|
| Compound | Pre diapause | Post diapause | Adult fly | ISTD |
| name | on column (ng) | on column (ng) | on column (ng) | on column (ng) |
| Histidine | 20 | 50 | 50 | 1 |
| Serine | 8 | 80 | 80 | 1 |
| Histamine | 3.2 | 3.2 | 3.2 | 1 |
| Aspartate | 25 | 250 | 250 | 1 |
| Glutamate | 25 | 250 | 250 | 1 |
| GABA | 10 | 100 | 100 | 1 |
| Dopa | 0.16 | 0.16 | 0.1 | 1 |
| Octopamine | 1 | 1 | 1 | 1 |
| Tyrosine | 4 | 40 | 40 | 1 |
| Dopamine | 1 | 1 | 1 | 1 |
| Tyramine | 0.32 | 0.32 | 0.64 | 1 |
| Serotonin | 1 | 1 | 1.6 | 1 |
| Tryptophan | 1 | 8 | 8 | 1 |
| Tryptamine | 1 | 1 | 0.16 | 1 |

*The 14 neurotransmitters were weighed individually and standard stock solutions of 1mg/ml in 0.1N HCL were prepared. The final concentration for each chemical for each experiment was calculated by running a preliminary experiment with calibration curves used for honey bee whole brain as in Ramesh et al 2019 and three replicates of apple/ hawthorn brain samples. The internal standard (ISTD) stock solution of 1mg/ml in 0.1NHCL was also prepared.